

10/511556
DT04 Rec'd PCT/PTO 15 OCT 2004

SEQUENCE LISTING

<110> Bayer AG

<120> Regulation of human transient receptor potential channel

<130> RCK-27

<140>

<141>

<150> US60/372899

<151> 2002-04-16

<150> US60/375139

<151> 2002-04-22

<160> 21

<170> PatentIn Ver. 2.1

<210> 1

<211> 5668

<212> DNA

<213> Homo sapiens

<400> 1

```
gtcacttagg aaaaggtgtc ctttcgggca gccgggctca gcatgaggaa cagaaggaat 60
gacactctgg acagcacccg gaccctgtac tccagcgcgt ctcggagcac agacttgtct 120
tacagtga aa ggcacttggg gaattttatt caagcaaatt ttaagaaacg agaattgtgc 180
ttctttacca aagattccaa ggccacggag aatgtgtgca agtgtggcta tgcccagagc 240
cagcacatgg aaggcaccca gatcaaccaa agtgagaaat ggaactacaa gaaacacacc 300
aaggaaatctc ctaccgacgc ctttggggat attcagtttg agacactggg gaagaaaggg 360
aagtatatat gtcgtgtcctg cgacacggac gcggaaatcc ttacgagct gctgacccag 420
cactgggcacc tgaaaacacc caacctgggc atttctgtga ccggggggcg caagaacttc 480
gccctgaagc cgcgcacatgc caagatcttc agcgggctca tctacatcgc gcagtccaaa 540
gggtgcttga ttctcacggg aggcacccat tatggcctga cgaagtacat cggggagggtg 600
gtgagagata acaccatcag caggagttca gaggagaata ttgtggccat tggcatagca 660
gcttggggca tgggtctccaa ccgggacacc ctcatcagga attgcatgct tgagggtctat 720
tttttagccc agtaccttat ggatgacttc acaagggatc cactgtatat cctggacaac 780
aaccacacac atttgctgct cgtggacaat ggctgtcatg gacatcccac tgtcgaagca 840
aagctccgga atcagctaga gaagcatatc tctgagcgca ctattcaaga ttccaactat 900
gggtggcaaga tccccattgt gtgttttgcc caaggagggtg gaaaagagac tttgaaagcc 960
atcaatacct ccatcaaaaa taaaattcct tgtgtggtgg tggaaggctc gggccggatc 1020
gctgatgtga tcgctagcct ggtggagggtg gaggatgccc cgacatcttc tgccgtcaag 1080
gagaagctgg tgcgttttt accccgcacg gtgtcccggt tgtctgagga ggagactgag 1140
agttggatca aatggctcaa agaaattctc gaatgttctc acctattaac agttattaaa 1200
atggaagaag ctggggatga aattgtgagc aatgccatct cctacgctct atacaaagcc 1260
ttcagcacca gtgagcaaga caaggataac tggaatgggc agctgaagct tctgctggag 1320
tggaaccagc tggacttagc caatgatgag attttcacca atgaccgccc atgggagctc 1380
gctgaccttc aagaagtcac gtttacggct ctcataaagg acagacccaa gtttgtccgc 1440
ctctttctgg agaattggctt gaacctacgg aagtttctca cccatgatgt cctcactgaa 1500
ctcttctcca accacttcag cagccttggt taccggaatc tgcagatcgc caagaattcc 1560
tataatgatg ccctcctcac gtttgtctgg aaactgggtg cgaacttccg aagaggcttc 1620
cggaaggaaag acagaaatgg ccgggacgag atggacatag aactccacga cgtgtctcct 1680
attactcggc accccctgca agctctcttc atctgggcca ttcttcagaa taagaaggaa 1740
ctctccaaag tcatttgagg gcagaccagg ggctgcactc tggcagccct gggagccagc 1800
aagcttctga agactctggc caaagtgaag aacgacatca atgctgctgg ggagtcgag 1860
gagctggcta atgagtacga gaccggggt tctgagctgt tctactgagtg ttacagcagc 1920
gatgaagact tggcagaaca gctgctgggc tattctgtg aagcttgggg tggaagcaac 1980
tgtctggagc tggcggtgga ggccacagac cagcatttca ccgcccagcc tgggggtccag 2040
aatcttcttt ctaagcaatg gtatggagag atttcccag acaccaagaa ctggaagatt 2100
atcctgtgtc tgtttattat acccttggtg ggtgtgggt ttgtatcatt taggaagaaa 2160
cctgtcgaca agcacaagaa gctgctttgg tactatgtgg cgttcttcac ctcccccttc 2220
gtggtcttct cctggaatgt ggtcttctac atcgcttcc tcctgctgtt tgcctacgtg 2280
ctgctcatgg atttccattc ggtgccacac cccccgagc tggctcctga ctgcgtggtc 2340
tttgcctctc tctgtgatga agtgagacag tggtaacgtaa atgggggtgaa ttattttact 2400
```

```

gacctgtgga atgtgatgga cacgctgggg cttttttact tcatagcagg aattgtatatt 2460
cggctccact cttctaataa aagctctttg tattctggag gaggcatttt ctgtctggac 2520
tacattatatt tcactctaag attgatccac atttttactg taagcagaaa cttaggaccc 2580
aagattataa tgctgcagag gatgctgac gatgtgttct tcttctgtt cctctttgcg 2640
gtgtggatgg tggcctttgg cgtggccagg caagggatcc ttaggcagaa tgagcagcgc 2700
tggaggtgga tattccgttc ggtcatctac gagccctacc tggccatggt cggccagggtg 2760
cccagtgaag tggatggtac cagtatgac tttgccact gcaccttcac tgggaatgag 2820
tccaaagcac tgggtgtgga gctggatgag cacaacctgc cccggttccc cgagtggatc 2880
accatcccc tgggtgtgcat ctacatgta tccaccaaca tctgtctggt caacctgctg 2940
gtcgccatgt ttggctacac ggtgggcacc gtccaggaga acaatgacca ggtctggaag 3000
ttccagaggt acttctctgg gcaggagtac tgcagcgcgc tcaatatccc cttccccttc 3060
atcgtcttcg cttacttcta catggtgttg aagaagtgt tcaagtgttg ctgcaaggag 3120
aaaaacatgg agtcttctgt ctgctgtttc aaaaatgaag acaatgagac tctggcatgg 3180
gaggggtgtca tgaaggaaaa ctacctgttc aagatcaaca caaaagccaa cgacacctca 3240
gaggaatga ggcctcgtat tagacaactg gatacaaac ttaatgatct caagggctct 3300
ctgaaagaga ttgctaataa aatcaataa actgtatga aactctaatt gagaataatc 3360
taattatagc aagatcatat taaggaatgc tgatgaacaa ttttgctatc gactactaaa 3420
tgagaatatt tcagaccctc ggttacatgg tggatgattt taaatcacc tagtgtgctg 3480
agaccttgag aataaagtgt gtgattgggt tcactctga agacggatat aaaggaagaa 3540
tatttctctt atgtgtttct ccagaatggg gctgtttct ctctgtgtct caatgcctgg 3600
gactggaggt tgatagttta agtgtgttct taccgctcc ttttctctt aatcttattt 3660
ttgatgaaca catatatagg agaacatcta tcttatgaat aagaacctgg tcatgcttta 3720
ctcctgtatt gttattttgt tcatttccaa ttgattctct acttttccct tttttgtatt 3780
atgtgactaa ttagtggca tttgttaaa agtctctcaa attaggccag attctaaaac 3840
atgtgcagc aagaggaccc cgctctctc aggaaaagt ttttcatttc tcaggatgct 3900
tcttacctgt cagaggaggt gacaaggcag tctctgtct tcttggactc accaggctoc 3960
tattgaagga accaccccca ttcctaataa tgtgaaaagt cgcccaaaat gcaaccttga 4020
aaggcactac tgactttgtt cttattggat actcctcta tttattattt ttcattataa 4080
aataatagct ggctattata gaaaatttag acatacaga gatgtagaaa gaacataaat 4140
tgtcccatc accttaaggt aatcactgct aacaatttct ggatggtttt tcaagtctat 4200
ttttttctta tgtatgtctc aattctcttt caaaatttta cagaatgtta tcatactaca 4260
tatatacttt ttatgtaagc tttttcactt agtattttat caaatatggt tttattatat 4320
tcatagcctt cttaaacatt atatcaataa ttgcataata ggcaacctct agcgattacc 4380
ataattttgc tcattgaagg ctatctccag ttgatcattg ggatgagcat ctttgtgcat 4440
gaatcctatt gctgtatttt ggaaaatttt ccaaggttag attccaataa atatctattt 4500
attattaaat attaaaatat cgatttatta ttaaaacat ttataaggct tttcataaaa 4560
tgtatgcaa ataggaaatta ttaacttgag cataagatat gagatacatg aacctgaact 4620
attaaaataa aatattatat ttaaccctag tttaagaaga agtcaatatg cttatttaaa 4680
tattatggat ggtgggcaga tcacttgagg tcaggagttc gagaccagcc tggccaacat 4740
ggcaaaacca catctctact aaaaataaaa aaattagctg ggtgtggtgg tgcactcctg 4800
taatcccagc tactcagaag gctgaggtac aagaattgct ggaacctggg aggcggaggt 4860
tgcaagtgaac caagattgca ccactgcact ccagccgggg tgacagagtg agactccgac 4920
tgaaaataaa taaataaata aataaataaa taaataaata aatattatgg atggtgaagg 4980
gaatggtata gaattggaga gattatctta ctgaacacct gtagtcccag ctttctctgg 5040
aagtgtggt atttgagcag gatgtgcaca aggcatttga aatgcccata attagttct 5100
cagctttgga tacactataa actcagtggtc tgaaggagga aatttttaga ggaagctact 5160
aaaagatcta atttgaaaaa ctacaaaagc atttactaaa aaagtttatt ttccttttgt 5220
ctgggcagta gtgaaaataa ctactcacia cattcactat gtttgcaagg aattaacaca 5280
aataaaagat gcctttttac ttaaacgcca agacagaaaa cttgcccaat actgagaagc 5340
aacttgcat agagagggaa ctgttaaatg ttttcaaccc agttcatctg gtggatgttt 5400
ttgcagggtta ctctgagaat tttgcttatg aaaaatcatt atttttagtg tagttcacaa 5460
taatgtattg aacatacttc taatcaaagg tgctatgtcc ttgtgtatgg tactaaatgt 5520
gtcctgtgta cttttgcaca actgagaatc ctgcggcttg gtttaatgag tgtgttcag 5580
aaataaataa tggaggaatt gtcaaaaaaa aaaaaaaa aaaaaaaa 5640
aaaaaaaa aaaaaaaaa aaaaaaaa 5668

```

<210> 2

<211> 3639

<212> DNA

<213> Homo sapiens

<400> 2

```

gattacgcaa gctatttagg tgacactata gaatwctcag cttgcatcaa gcttgggtacc 60
gagctcggat ccctagtaac ggccgccagt gtgctggaat tcgcccttgc agccgggctc 120
agcatgagga ccagaaggaa tgacactctg gcagcaccc ggacctgtga ctccagcgcg 180
tctcggagca cagactgtgc ttacagtga agcgacttgg tgaattttat tcaagcaaat 240
tttaagaaac gagaatgtgt cttctttacc aaagattcca aggccacgga gaatgtgtgc 300

```

```

aagtgtggct atgccagag ccagcacatg gaaggcacc agatcaacca aagtgagaaa 360
tggaaactaca agaaacacac caaggaatgt cctaccgacg cctttgggga tattcagttt 420
gagacactgg ggaagaaagg gaagtatata cgtctgtcct gcgacacgga cgcggaaatc 480
ctttacgagc tgctgaccca gcaactggcag ctgaaaacac ccaacctggg catttctgtg 540
accggggggc ccaagaactt cgcctgaag ccgcgcagtc gcaagatctt cagccggctc 600
atctacatcg cgcagtccaa aggtgcttgg attctcacgg gaggcacca ttatggcctg 660
atgaagtaca tcggggaggt ggtgagagat aacaccatca gcaggagttc agaggagaat 720
attgtggcca ttggcatagc agcttggggc atgggtctcca accgggacac cctcatcagg 780
aattgcatg ctgaggggcta ttttttagcc cagtacctta tggatgactt cacaagagat 840
ccactgtata tcctggacaa caaccacaca catttgtgc tcgtggacaa tggctgtcat 900
ggacatccca ctgtcgaagc aaagctccgg aatcagctag agaagtatat ctctgagcgc 960
actattcaag attccaacta tgggtggcaag atccccattg tgtgttttgc ccaaggaggt 1020
ggaaaagaga ctttgaaagc catcaatacc tccatcaaaa ataaaaattc ttgtgtgtg 1080
gtggaaggct cggggcagat cgctgatgtg attcgtagcc tgggtggagg ggaggatgcc 1140
ctgacatctt ctgcccgtcaa ggagaagctg gtgcgctttt taccgcgac ggtgtcccgc 1200
ctgctgagg aggagactga gaggttggatc aaatggctca aagaaattct cgaatgttct 1260
cacctattaa cagttattaa aatggaagaa gctggggatg aaattgtgag caatgccatc 1320
tcctacgctc tatacaaagc cttcagcacc agtgagcaag acaaggataa ctggaattgg 1380
cagctgaagc ttctgtgga gtggaaccag ctggacttag ccaatgatga gattttcacc 1440
aatgaccgcc gatgggagtc tgctgacctt caagaagtca tgtttacggc tctcataaag 1500
gacagaccca agtttgtccg cctctttctg gagaatggct tgaacctacg gaagtttctc 1560
accatgatg tcctcactga actcttctcc accacttca gcacgcttgt gtaccggaat 1620
ctgcagatcg ccaagaattc ctataatgat gctctctca cgtttgtctg gaaactgggt 1680
gcgaacttcc gaagaggctt ccggaaggaa gacagaaatg gccgggacga gatggacata 1740
gaactccacg acgtgtctcc tattactcgg caccctctgc aagctctctt catctgggcc 1800
attcttcaga atagaagga actctccaaa gtcatttggg agcagaccag gggctgcact 1860
ctggcagccc tgggagccag caagcttctg aagactctgg ccaaagtga gaacgacatc 1920
aatgtgctg gggagtcgga ggagctggct aatgagtacg agaccgggc tgttgagctg 1980
ttcactgagt gttacagcag cgatgaagac ttggcagaac agctgctggt ctattcctgt 2040
gaagcttggg gtggaagcaa ctgtctggag ctggcggtgg aggccacaga ccagcatttc 2100
atgcctgagc atggggctca gaattttctt tctaagcaat ggtatggaga cattttcact 2160
gacaccaaga actggaagat tatcctgtgt ctgtttatta tacccttggg gggctgtggc 2220
tttgtatcat ttaggaagaa acctgtcgac aagcacaaga agctgctttg gtactatgtg 2280
gcgttcttca cctccccctt cgtggtcttc tcttggaaat tggctcttca catcgccttc 2340
ctcctctgt tgcctacgt gctgctcatg tacttccatt cggtgccaca ccccccgag 2400
ctggctctgt actcgtggt ctttgtcctt ttctgtgatg aagtgagaca gtggtacgta 2460
aatggggtga attattttac tgacctgtgg aatgtgatgg acacgctggg gcttttttac 2520
ttcatagcag gaattgtatt tcggtctccac tcttctaata aaagctcttt gtattctgga 2580
cgagtcattt ttgtctgga ctacattatt ttcactctaa gattgatcca catttttact 2640
gtaagcagaa acttaggacc caagattata atgctgcaga ggatgctgat cgatgtgttc 2700
ttcttctgt tcctctttgc gwggtggatg gtggcctttg gcgtggccag gcaagggatc 2760
cttaggcaga atgagcagcg ctggaggtgg atattccgtt cggtcatcta cgagccctac 2820
ctggccatgt tcggccaggt gccagtgac gtggatggtg ccacgtatga ctttggccac 2880
tgcacattca ctgggaatga cccaagcca gtctgtgtgg agctggatga gcacaacctg 2940
ccccggttcc ccgagtggat caccatcccc ctggtgtgca tctacatgtt atccaccaac 3000
atcctgctgg tcaacctgct ggtcgccatg tttggctaca cgggtgggac cgtccaggag 3060
aacaatgacc aggtctggaa gttccagagg tacttctgg tgcaggagta ctgcagccgc 3120
ctcaatatcc ccttccccct catcgtcttc gcttacttct acatgggtgg gaagaagtgc 3180
ttcaagtgtt gctgcaagga gaaaacatg gagtctctg tctgctgttt caaaaatgaa 3240
gacaatgaga ctctggcatg ggaggggtgc atgaaggaaa actacctgt caagatcaac 3300
acaaaagcca acgacacctc agaggaaatg aggcacgat ttagacaact ggatacaaag 3360
cttaatgatc tcaaggggtc tctgaaagag attgttaata aaactgtagt aaactgtagt 3420
aactctaag gagaaaaatc taattatagc aagatcatat taaggaaatgc tgatgaacaa 3480
ttttgctatc gactactaaa tgagagattt tcagaccctt gggtagatgg tggatgattt 3540
taaatacccc tagtgtgtg agaccttgag aataaagtgt gaagggcgaa ttctgcagat 3600
atccatcaca ctggcggccg ctcgagcatg catctagag 3639

```

<210> 3

<211> 3042

<212> DNA

<213> Homo sapiens

<400> 3

```

atggttggag gatgcagggt gacagaagac gtggagcctg cagaagtaaa ggaaaagatg 60
tcctttcggg cagccagggt cagcatgagg aacagaagga atgacactct ggacagcacc 120
cggaccctgt actccagcgc gtctcggagc acagacttgt cttacagtga aagcgacttg 180
gtgaatttta ttcaagcaaa ttttaagaaa cgagaatgtg tcttctttac caaagattcc 240

```

```

aaggccacgg agaattgtgtg caagtgtggc tatgcccgaga gccagcacat ggaaggcacc 300
cagatcaacc aaagtgaagaa atggaaactac aagaaacaca ccaaggaatt tcctaccgac 360
gcctttgggg atattcagtt tgagacactg ggaagaaag ggaagtatat acgtctgtcc 420
tgcgacacgg acgcggaaat cctttacgag ctgctgaccc agcactggca cctgaaaaca 480
cccaacctgg tcatttctgt gaccgggggc gccagaact tcgccctgaa gccgcgcatg 540
cgcaagatct tcagccggct catctacatc gcgcagtcca aaggtgcttg gattctcacg 600
ggaggcacc attatggcct gatgaagta ctcggggagg tggtagaga taacaccatc 660
agcaggagtt cagaggagaa tattgtggcc attggcatag cagcttgggg catggtctcc 720
aaccgggaca ccctcatcag gaattgcgat gctgagggtc attttttagc ccagtacctt 780
atggatgact tcacaagaga tccactgtat atcctggaca acaaccacac acatttgctg 840
ctcgtggaca atggctgtca tggacatccc actgtcgaag caaagctccg gaatcgacta 900
gagaagtata tctctgagcg cactattcaa gattccaact atgggtggca gatccccatt 960
gtgtgttttg cccaaggagg tggaaaagag actttgaaag ccatcaatac ctccatcaaa 1020
aataaaattc cttgtgtggt ggtggaaggc tcgggcccaga tcgctgatgt gatcgctagc 1080
ctggtggagg tggaggatgc cctgacatct tctgccgta aggagaagct ggtgcgcttt 1140
ttaccccgca cgggtgtccc gctgcctgag gaggagactg agagttggat caaatggctc 1200
aaagaaattc tcgaatgttc tcacctatta acagttatta aaatggaaga agctggggat 1260
gaaattgtga gcaatgccat ctctacgct ctatacaaag ccttcagcac cagttagcaa 1320
gacaaggata actggaatgg gcagctgaag cttctgtg agtggaaacca gctggactta 1380
gccaatgatg agattttcac caatgaccgc cgatgggaga agagcaaacc gaggtcaga 1440
gacacaataa tccaggtcac atggctggaa aatggtagaa tcaaggttga gagcaaagat 1500
gtgactgacg gcaaagcctc ttctcatatg ctggtggttc tcaagtctgc tgaccttcaa 1560
gaagtcatgt ttacggctct cataaaggac agaccgaagt ttgtccgctt cttctggag 1620
aatggcttga acctacggaa gtttctcacc catgatgtcc tcaactgaact cttctccaac 1680
cacttcagca cgcttgtgta ccggaatctg cagatcgcca agaattccta taatgatgcc 1740
ctcctcacgt ttgtctggaa actggttgcg aacttccgaa gaggttccg gaaggagac 1800
agaaatggcc gggacgagat ggacatagaa ctccacgacg tgtctcctat tactcggcac 1860
cccctgcaag ctctcttcac ctgggcccatt ttccagaata agaaggaaact ctccaaagtc 1920
atltgggagc agaccagggg ctgcactctg gcagccctgg gagccagcaa gcttctgaag 1980
actctggcca aagtgaagaa cgacatcaat gctgctgggg agtccgagga gctggctaata 2040
gagtagctga cccgggctgt tggtagtccc acagtgtgga atgctgtggt gggcgcggaat 2100
ctgcatgtg gcacagacat tgccagcggc actcatagac cagatgggtg agagctgttc 2160
actgagtgtt acagcagcga tgaagacttg gcagaacagc tgctggtcta ttctgtgaa 2220
gcttgggggt gaagcaactg tctggagctg gcggtggagg ccacagacca gcatttcatc 2280
gcccagcctg ggggtccagaa ttttctttct aagcaatggt atggagagat ttcccagac 2340
accaagaatt ggaagaaacc tctgtgtctg cacaagaagc tgctttggta ctatgtggcg 2400
gtatcattta ccccttctgt ggtcttctcc ttggaatgtg tcttctacat cgccttctcc 2520
ctgctgtttg cctacgtgct gctcatggat ttccattcgg tgccacaccc ccccgagctg 2580
gtcctgtact cgctggtctt tgtcctcttc tgcagtaag tgagacaggg ccgcccggct 2640
gctcccagtg cggggcccgc caagcccacg cccaccggga actccatctg gcccgcaagc 2700
tccacacgca gcccgggttc ccgctcacgc cactccttcc acacttccct gcaagctgag 2760
ggtgccagct ctggccttgg ccagcccaga aaggggctcc cacagtgcag cgggtgggctg 2820
aagggctcct caagtgcgcg caaagtggga gccccgcag aggaggtgcc gagagcaagc 2880
gagggtctgt aggactgcca gcaagctgtc acctctcaga agcgtaagac agcaatggac 2940
caaacagacg aagatctctt cccctatgga gcattctacc agttcctgat gatttccagg 3000
agctttcgag gagaggagat gagcatcggc aagcagcact aa 3042

```

<210> 4

<211> 3039

<212> DNA

<213> Homo sapiens

<400> 4

```

atggttgagg gatgcagggt gacagaagac gtggagcctg cagaagtaaa ggaaaagatg 60
tcctttcggg cagccaggct cagcatgagg aacagaagga atgacactct ggacagcacc 120
cggaccctgt actccagcgc gtctcggagc acagacttgt cttacagtga aagcgacttg 180
gtgaatttta ttcaagcaaa ttttaagaaa cgagaatgtg tcttctttac caaagattcc 240
aaggccacgg agaattgtgt caagtgtggc tatgcccgaga gccagcacat ggaaggcacc 300
cagatcaacc aaagtgaagaa atggaactac aagaaacaca ccaaggaatt tcctaccgac 360
gcctttgggg atattcagtt tgagacactg ggaagaaag ggaagtatat acgtctgtcc 420
tgcgacacgg acgcggaaat cctttacgag ctgctgaccc agcactggca cctgaaaaca 480
cccaacctgg tcatttctgt gaccgggggc gccagaact tcgccctgaa gccgcgcatg 540
cgcaagatct tcagccggct catctacatc gcgcagtcca aaggtgcttg gattctcacg 600
ggaggcacc attatggcct gatgaagtac ctcggggagg tggtagaga taacaccatc 660
agcaggagtt cagaggagaa tattgtggcc attggcatag cagcttgggg catggtctcc 720
aaccgggaca ccctcatcag gaattgcgat gctgagggtc attttttagc ccagtacctt 780

```

- 5 -

```

atggatgact tcacaagaga tccactgtat atcctggaca acaaccacac acatttgctg 840
ctcgtggaca atggctgtca tggacatccc actgtcgaag caaagctccg gaatcagcta 900
gagaagtata tctctgagcg cactattcaa gattccaact atggtggcaa gatccccatt 960
gtgtgttttg cccaaggagg tggaaaagag actttgaaag ccatcaatac ctccatcaaa 1020
aataaaattc cttgtgtggg ggtggaaggc tcggggccaga tcgctgatgt gatcgctagc 1080
ctggtggagg tggaggatgc cctgacatct tctgccgtca aggagaagct ggtgcgcttt 1140
ttaccccgca agtgtgtccc gctgcctgag gaggagactg agagtggat caaatggctc 1200
aaagaaattc tcgaatgttc tcacctatta acagttatta aaatggaaga agctggggat 1260
gaaattgtga gcaatgccat ctctacgct ctatacaaag ccttcagcac cagtgaagaa 1320
gacaaggata actggaatgg gcagctgaag ctctgtctgg agtggaaacca gctggactta 1380
gccaatgatg agattttcac caatgaccgc cgttgggaga agagcaaacc gaggctcaga 1440
gacacaataa tccaggtcac atggctggaa aatggtagaa tcaaggttga gagcaaatgat 1500
gtgactgacg gcaaaagcctc ttctcatatg ctggtggttc tcaagtctgc tgacctcaa 1560
gaagtcatgt ttacggctct cataaaggac agaccaagt ttgtccgctt ctttctggag 1620
aatggcttga acctacggaa gtttctcacc catgatgtcc tcaactgaact cttctcgaac 1680
cacttcagca cgcttgtgta ccggaatctg cagatcgcca agaattccta taatgatgcc 1740
ctcctcacgt ttgtctggaa actggttgcg aacttccgaa gaggcttccg gaaggagac 1800
agaaatggcc gggacgagat ggacatagaa ctccacgacg tctctcctat tactcggcac 1860
ccctggaag ctctcttcat ctgggacatt cttcagaata agaaggaact ctccaaagtc 1920
atttgggagc agaccagggg ctgcactctg gcagccctgg gagccagcaa gcttctgaag 1980
actctggcca aagtgaagaa cgacatcaat gctgctgggg agtccgagga gctggctaatt 2040
gagtacctga cccgggctgt tggtagctcc acagtgtgga atgctgtggt gggcgcggaat 2100
ctgcatgtgt gcacagacat tgccagcggc actcatagac cagatggtgg agagtgttc 2160
actgagtgtt acagcagcga tgaagacttg gcagaacagc tgctggtcta ttctgtgaa 2220
gcttgggggtg gaagcaactg tctggagctg gcggtggagg ccacagacca gcatttcac 2280
gccacgcctg ggggtccagaa ttttctttct aagcaatggt atggagagat ttcccagac 2340
accaagaact ggaagattat cctgtgtctt tttattatac ccttgggtgg ctgtggttc 2400
gtatcattta ggaagaaacc tgtcgacaag cacaagaagc tgctttggtg ctatgtggcg 2460
ttcttcacct ccccttctgt ggtcttctcc tggaaatgtg tcttctacat cgcttctc 2520
ctgctgtttg cctacgtgct gctcatggat ttccattcgg tgccacaccc ccccgagctg 2580
gtctgttact cgctggtctt tgtcctcttc tbtgatgaag tgagacaggg ccggcgggct 2640
gctccagtg cggggccgc caagcccacg gcccacccgga actccatctg gcccgcaagc 2700
tccacacgca gccccgggtc ccgctcacgc cactccttcc acacttccct gcaagctgag 2760
ggtgccagct ctggccttgg ccagcccaga aaggggctcc cacagtgcag cgggtgggctg 2820
aagggctcct caagtgcgcg caaagtggga gccacggcag aggaggtgcc gagagcaagc 2880
gagggctgtg gcagctgcca acctctcaga agcgtaagac agcaatggac 2940
caaacagacg aagatctctt cccctatgga gcattctacc agttcctgat gatttccagg 3000
agctttcgag gagaggagat gagcatcggc aagcagcac 3039

```

<210> 5

<211> 3893

<212> DNA

<213> Homo sapiens

<400> 5

```

atataagttt ataaaaacagt ggctggatgg ttggaggatg caggtggaca gaagacgtgg 60
agcctgcaga agtaaaaggaa aagatgtcct ttcgggcagc caggctcagc atgaggaaca 120
gaaggaatga cactctggac agcaccgcga ccctgtactc cagcgcgtct cggagcacag 180
acttgtctta cagtgaagac gccagcttct acgctgcctt caggacacag acgtgcccac 240
tcatggcttc ttgggacttg gtgaatttta ttcaagcaaa ttttaagaaa cgagaatgtg 300
tcttctttac caaagattcc aaggccacgg agaattgtgt caagtgtggc tatgccaga 360
gccagcacat ggaaggcacc cagatcaacc aaagtgaaga atggaactac aagaaacaca 420
ccaaggaatt tcctaccgac gcctttgggg atattcagtt tgagacactg ggaagaaag 480
ggaagtatat acgtctgtcc tgcgacacgg acgcggaaat cctttacgag ctgctgacct 540
agcactggca cctgaaaaa cccaacctgg tcatttctgt gaccgggggc gccagaaact 600
tcgcccgtga gccgcgcatg cgcaagatct tcagccggct catctacatc gcgcagtcca 660
aaggtgcttg gattctcacg ggaggcacc attatggcct gatgaagta atcggggagg 720
tggtgagaga taacaccatc agcaggagtt cagaggagaa tattgtggcc attggcatag 780
cagcttgggg catggtctcc aaccgggaca ccctcatcag gaattgcgat gctgagggct 840
attttttagc ccagtacctt atggatgact tcacaagaga tccactgtat atcctggaca 900
acaaccacac acatttgctg ctctgtggaca atggctgtca tggacatccc actgtcgaag 960
caaagctccg gaatcagcta gagaagtata tctctgagcg cactattcaa gattccaact 1020
atggtggcaa gatccccatt gtgtgttttg cccaaggagg tggaaaagag actttgaaag 1080
ccatcaataa ctccatcaaa aataaaattc cttgtgtggg ggtggaaggc tcggggccaga 1140
tcgctgatgt gatcgctagc ctggtggagg tggaggatgc cctgacatct tctgccgtca 1200
aggagaagct ggtgcgcttt ttaccccgca cgggtgtccc gctgcctgag gaggagactg 1260
agagttggat caaatggctc aaagaaattc tcgaatgttc tcacctatta acagttatta 1320

```

| | | | | | | |
|-------------|-------------|-------------|-------------|------------|-------------|------|
| aaatggaaga | agctggggat | gaaattgtga | gcaatgccat | ctcctacgct | ctatacaaa | 1380 |
| ccttcagcac | cagtgagcaa | gacaaggata | actggaatgg | gcagctgaag | cttctgctgg | 1440 |
| agtggaaacca | gctggactta | gccaatgatg | agattttcac | caatgaccgc | cgatgggaga | 1500 |
| agagcaaacc | gaggctcaga | gacacaataa | tccaggtcac | atggctggaa | aatggtagaa | 1560 |
| tcaaggttga | gagcaaagat | gtgactgacg | gcaaagcctc | ttctcatatg | ctggtgggttc | 1620 |
| tcaagctcgc | tgacctcaa | gaagtcattg | ttacggctct | cataaaggac | agaccctaag | 1680 |
| ttgtccgcct | ctttctggag | aatggcttga | acctacggaa | gtttctcacc | catgatgtcc | 1740 |
| tcactgaact | cttctccaac | cacttcagca | cgcttgtgta | ccggaatctg | cagatcgcca | 1800 |
| agaattccta | taatgatgcc | ctcctcacgt | ttgtctggaa | actggttgcg | aacttccgaa | 1860 |
| gaggcttccg | gaaggaagac | agaaatggcc | gggacgagat | ggacatagaa | ctccacgacg | 1920 |
| tgtctcctat | tactcggcac | cccctgcaag | ctctcttcat | ctgggccatt | cttcagaata | 1980 |
| agaaggaact | ctccaaagtc | atttgggagc | agaccagggg | ctgcactctg | gcagccctgg | 2040 |
| gagccagcaa | gcttctgaag | actctggcca | aagtgaagaa | cgacatcaat | gctgtgggg | 2100 |
| agtccgagga | gctggctaata | gagtagcaga | cccgggctgt | tggtgagtc | acagtgtgga | 2160 |
| atgctgtggt | gggcgcggat | ctggcatgtg | gcacagacat | tgccagcggc | actcatagac | 2220 |
| cagatgggtg | agagctgttc | actgagtgtt | acagcagcga | tgaagacttg | gcagaacagc | 2280 |
| tgctggtcta | ttcctgtgaa | gcttgggggtg | gaagcaactg | tctggagctg | gcggtggagg | 2340 |
| ccacagacca | gcatttcata | gcccagcctg | gggtccagaa | ttttctttct | aagcaatggt | 2400 |
| atggagagat | tcccagagac | accaagaatt | ggaagattat | cctgtgtctg | tttattatac | 2460 |
| ccttgggtggg | ctgtggcttt | gtatcattta | ggagaaacc | tgctgacaag | cacaagaagc | 2520 |
| tgcttggta | ctatgtggcg | ttcttcacct | cccccttctg | ggctcttctc | tggaatgtgg | 2580 |
| tcttctacat | cgcttctctc | ctgtgttttg | cctacgtgct | gctcatggat | ttccattcgg | 2640 |
| tgccacaccc | ccccgagctg | gtcctgtact | cgctggctct | tgctctcttc | tgtatgaag | 2700 |
| tgagacaggg | ccggccggct | gctcccagtg | cggggcccg | caagcccacg | cccacccgga | 2760 |
| actccatctg | gcccgcagc | tcacacgcga | gccccgggtc | ccgctcacgc | cactccttcc | 2820 |
| acacttccct | gcaagctgag | ggtgccagct | ctggccttgg | ccagcccaga | aagggggtgga | 2880 |
| catttaaaaa | tctggaaatg | gttgatattt | ccaagctgct | gatgtccctc | tctgtccctt | 2940 |
| tctgtacgca | gtggtagta | aatggggtga | attattttac | tgacctgtgg | aatgtgatgg | 3000 |
| acacgctggg | gcttttttac | ttcatagcag | gaattgtatt | tcggcaagg | atccttaggc | 3060 |
| agaatgagca | gcgctggagg | tggtatttcc | gttcggctcat | ctacgagccc | tacctggcca | 3120 |
| tgctcggcca | tggtgccagt | gacgtggatg | gtaccacgta | tgactttg | cactgcacct | 3180 |
| tcactcggaa | gggtccaaag | ccactgtgtg | tggagctgga | tgagcacaac | ctgcgccggt | 3240 |
| tccccgagtg | gatcaccatc | ccccgtgtgt | gcactctacat | gttatccacc | aacatcctgc | 3300 |
| tggtcaacct | gctggctgcc | atgtttggct | acacgggtgg | cacogtccag | gagaaacaatg | 3360 |
| accaggtctg | gaagtccag | aggtacttcc | tggtgcagga | gtactgcagc | cgctcaata | 3420 |
| tccccttccc | cttcactgtc | ttcgcttact | tctacatggt | ggtgaagaag | tgcttcaatg | 3480 |
| gttgctgcaa | ggagaaaaac | atggagtctt | ctgtctgctg | tgagtgggtt | atccatgtgt | 3540 |
| acttgggatc | agaagcagcg | attaatttca | gggaaggatg | cctgcatcca | gtgattggaa | 3600 |
| gctggacccc | aggctggctg | gtctggacat | ccacacgc | tctcacatgc | agtgcgggct | 3660 |
| ggccagagca | aggaggtctc | agtgtcaca | cacatagcag | ctgggttctc | gcaaaaagca | 3720 |
| gcaagtcaca | ggccacacca | gacagacgg | gtagagaatg | tgactctgct | tctgggtggg | 3780 |
| aaggacagcc | tgcccggtgg | gtggaagaat | ccgtggccct | gtttggccat | cggtggccctg | 3840 |
| tttggccacc | taccactcta | ggcatcactg | agctgaatgc | gccggctctc | tga | 3893 |

<210> 6

<211> 4646

<212> DNA

<213> Homo sapiens

<400> 6

| | | | | | | |
|------------|------------|-------------|-------------|-------------|-------------|------|
| tcgacccacg | cgctccgcca | cgcgctccgcc | cacgcgctccg | cccacgcgctc | cgccacgcg | 60 |
| tccgcccacg | cgctccgggt | gaaagmramy | cmvgcktsms | aaaaaccgctc | acttaggaaa | 120 |
| agatgtcctt | tcgggcagcc | aggtcagca | tgaggacag | aaggaaatgac | actctggaca | 180 |
| gcacccggac | cctgtactcc | agcgctctc | ggagcacaga | cttgtcttac | agtgaagcg | 240 |
| acttgggtga | ttttattcaa | gcaaatttta | agaaacgaga | atgtgtcttc | tttaccaaag | 300 |
| attccaaggc | ccggagaaat | gtgtgcaagt | gtggctatgc | ccagagccag | cacatggaag | 360 |
| gcacccagat | caaccaaagt | gagaaatgga | actacaagaa | acacaccaag | gaatttcta | 420 |
| ccgacgcctt | tggggatatt | cagtttgaga | cactggggaa | gaaagggag | tatatagtc | 480 |
| tgctcctgca | cacggacgcg | gaaatccttt | acgagctgct | gacccagcac | tggcacctga | 540 |
| aaacacccaa | cctggtcatt | tctgtgaccg | ggggcgccaa | gaacttcg | ctgaagccgc | 600 |
| gcatgcgcaa | gatcttcagc | cggtctatct | agcatcgcca | gtccaaagg | gcttgatttc | 660 |
| tcacgggagg | cacccattat | ggcctgatga | agtacatcgg | ggaggtgggtg | agagataaca | 720 |
| ccatcagcag | gagttcagag | gagaatattg | tggtccattgg | catagcagct | tggtggcatgg | 780 |
| tctccaaccg | ggacacccct | atcaggaatt | gcgatgtctga | gggtattttt | ttagccagct | 840 |
| accttatgga | tgacttcaca | agagatccac | tgtgtatcct | ggacaacaac | cacacacatt | 900 |
| tgctgctcgt | ggacaatggc | tgtcatggac | atcccactgt | cgaagcaaa | ctccggaatc | 960 |
| agctagagaa | gtatatctct | gagcgcacta | ttcaagattc | caactatggt | ggcaagatcc | 1020 |

```

ccattgtgtg ttttgcccaa ggaggtggaa aagagacttt gaaagccatc aatacctcca 1080
tcaaaaaataa aattccttgt gtggtggtgg aaggctcggg ccagatcgct gatgtgatcg 1140
ctagcctggt ggaggtggag gatgccctga catcttctgc cgtcaaggag aagctggtgc 1200
gctttttacc ccgcacggtg tcccggctgc ctgaggagga gactgagagt tggatcaaat 1260
ggctcaaaaga aattctcgaa tgttctcacc tattaacagt tattaanaatg gaagaagctg 1320
gggatgaaat tgtgagcaat gccatctcct acgctctata caaagccttc agcaccagtg 1380
agcaagacaa ggataactgg aatgggcagc agatcgccaa gctggagtgg aaccagctgg 1440
acttagccaa tgatgagatt ttcaccaatg accgccgatg ggagtctgct gaccttcaag 1500
aagtcatggt tacggctctc ataaaggaca gacccaagt tgtccgcctc tttctggaga 1560
atggcttgaa cctacggaag tttctcacc atgatgtcct cactgaactc tttccaacc 1620
acttcagcac gcttggtgac cggaatctgc agatcgccaa gaattcctat aatgatgcc 1680
tcctcacgtt tgtctggaaa ctggttgcca acttccgaag aggttccgg aaggaagaca 1740
gaaatggccg ggacgagatg gacatagaac tccacgacgt gtctcctatt actcggcacc 1800
ccctgcaagc tctcttcac tgggccattc ttcagaataa gaaggaactc tccaaagtca 1860
tttggggcaa gaccagggca tgcactctgg agcccgctgg agccagcaag cttctgaaga 1920
ctctggccaa agtgaagaac gacatcaatg ctgctgggga gtccgaggag ctggctaag 1980
agtacgagac ccgggctggt gagctgttca ctgagtgtta cagcagcgt gaagacttgg 2040
cagaacagct gctggtctat tcctgtgaag cttggggtgg aagcaactgt ctggagctgg 2100
cgggtggagg cagacaccag catttctgc cccagcctgg ggtccagaat tttctttcta 2160
agcaatggta tggagagatt tcccagaca ccaagaactg gaagattatc ctgtgtctgt 2220
ttattatacc cttggtgggc tgtggctttg tatcatctag gaagaaacct gtccgaagc 2280
acaagaagct gctttggtac tatgtggcgt tcttcacctc ccccttcgtg gtcttctcct 2340
ggaatgtggt cttctacac gccttctcc tgcgtttgct ctacgtgctg ctcttgatt 2400
tccattcggg gccacacccc cccgagctgg tctgtactc gctggtcttt gtctctctt 2460
gtgatgaagt gagacagtgg tacgtaaatg ggggtgaatta tttactgac ctgtggaatg 2520
tgatggacac gctggggctt ttttacttca tagcaggaat tgtatttcgg ctccactctt 2580
ctaataaaag ctctttgtat tctggacgag tcattttctg tctggactac attatttcta 2640
ctctaagatt tttactgtaa gcagaaactt cagaaaactt cttgcggtg tggatggtg 2700
tgcagaggat gctgatcgat gtgttcttct tctgttctc cttgcggtg tggatggtg 2760
cctttggcgt ggccaggcaa gggatcctta ggcagaatga gcagcgctgg aggtggatat 2820
tccgttcggg catctacgag ccctacctgg ccattgtcgg ccagggtgcc agtgactgtg 2880
atggtaccac atgtacttt gccactgca ccttactgg gaatgagtc aagccactgt 2940
gtgtggagct ggatgagcac aacctgcccc ggttccccga gtggatcacc atccccctgg 3000
tgtgcatcta catgttatcc accaacatcc tgcgtggtcaa cctgctggtc gccatgtttg 3060
gctacacggt gggcacccgtc caggagaaca atgaccaggt ctggaagttc cagaggtact 3120
tcttggtgca gtagtactgc agccgcctca atatcccctt ccccttcac gtcttcgctt 3180
acttctacat ggtggtgaag aagtgttca agtgttgctg caaggagaaa aacatggagt 3240
cttctgtctg ctgtttcaaa aatgaagaca atgagactct ggcatgggag ggtgtcatga 3300
aggaaaacta ccttgtcaag atcaacacaa aagccaacga cacctcagag gaaatgagc 3360
atcgatttag acaactggat acaagctta atgtactcaa ggttctctg aaagagattg 3420
ctaataaaat caaataaaac tgtatgaact ctaatggaga aaaaactaat tatagcaaga 3480
tcataattaag gaatgctgat gaacaatttt gctatcgact actaaatgag agattttcag 3540
acccttgggt acatgggtgga tgattttaaa tcaccctagt gtgctgagac cttgagaata 3600
aagtgtgtga ttggtttcat acttgaagac ggatataaag gaagaatatt tcccttatgt 3660
gtttctccag aatgggtgct gtttctctct gtgtctcaat gcctgggact ggaggttgat 3720
agttaagtgt tgttcttacc gcctcctttt tcctttaatc ttatttttga tgaacacata 3780
tataggagaa catctatcct atgaataaga acctggtcat gctttactcc tgtattgtta 3840
ttttgttcat ttccaattga ttctctactt ttcccttttt tgtattatgt gactaattag 3900
ttggcatatt gtwaaggatc tctcaaatta ggccagattc taaaacatgc tgcagcaaga 3960
ggaccccgct ctcttcagga aaagtgtttt catttctcag gatgcttctt acctgtcaga 4020
ggaggtgaca aggcagtctc ttgtctctct ggactacca ggctcctatt gaaggaacca 4080
ccccatttcc taaatatgtg aaaagtgcgc caaaatgcaa ccttgaaaag cactatgac 4140
tttgttctta ttggatactc ctcttattta ttatttttcc attaaaaata atagctggct 4200
attatagaaa atttagacca tacagagatg tagaaagaac ataaattgtc cccattacct 4260
taaggtaatc actgctaaca atttctggat ggtttttcaa gtctattttt tttctatgta 4320
tgtctcaatt ctctttcaaa attttacaga atgttatcat actacatata tacttttat 4380
gtaagctttt tcacttagta ttttatcaaa tatgttttta ttatattcat agccttctta 4440
aacattatat caataattgc ataataggca acctctagcg attaccataa ttttgctcat 4500
tgaaggctat ctccagttga tcattgggat gagcatcttt gtgcatgaat cctatttgctg 4560
tatttgggaa aattttccaa ggttagattc caataaatat ctatttatta ttaaaaaaaa 4620
aaaaaaaaag gcgccgctc tagagt 4646

```

<210> 7

<211> 3639

<212> DNA

<213> Homo sapiens

<400> 7

```

gattacgcaa gctattttagg tgacactata gaatwctcag cttgcatcaa gcttgggtacc 60
gagctcggat ccctagtaac ggccgccagtg gtgctggaat tcgcccttgc agccggggctc 120
agcatgagga acagaaggaa tgacactctg gacagcaccg ggaccctgta ctccagcgcg 180
tctcggagca cagacttgct ttacagtgaag agcagcttgg tgaattttat tcaagcaaat 240
tttaagaaac gagaatgtgt cttcttttacc aaagattcca agggccacgga gaatgtgtgc 300
aagtgtggct atgcccagag ccagcacatg gaaggcacc agatcaacca aagtgaagaaa 360
tggaactata agaaacacac caaggaatct cctaccgacg cctttgggga tattcagttt 420
gagacactgg ggaagaaagg gaagtatata cgtctgtcct gcgacacgga cgcggaaatc 480
ctttacgagc tgctgaccca gcactggcac ctgaaaacac ccaacctggt catttctgtg 540
accggggggcg ccaagaactt cgccctgaag ccgcgcacg gcaagatctt cagccggctc 600
atctacatcg cgcagtcctaa aggtgcttgg attctcacgg gaggcacca ttatggctcg 660
atgaagtaca tcggggaggt ggtgagagat aacaccatca gcaggagttc agaggagaat 720
attgtggcca ttggcatagc agcttggggc atggtctcca accgggacac cctcatcagg 780
aattgcgatg ctgagggcta ttttttagcc cagtacctta tggatgactt cacaagagat 840
ccactgtata tctggacaa caactgtgct cgttggacaa tggctgtcat 900
ggacatccca ctgtcgaagc aaagctccgg aatcagctag agaagtatat ctctgagcgc 960
actattcaag attocaacta tgggtggcaag atccccattg tgtgttttgc ccaaggagggt 1020
ggaaaagaga ctttgaaagc catcaatacc tccatcaaaa ataaaattcc ttgtgtgggt 1080
ctgaaggtcg cgggccagat cgctgatgtg attcgtacgg tgggtggagg ggaggtgccc 1140
ctgacatctt ctgccgtcaa ggagaagctg gtgcgctttt taccgccgac ggtgtcccgg 1200
ctgcctgagg aggagactga gagtgtgac aaatggctca aagaaattct cgaatgttct 1260
cacctattaa cagttattaa aatggaagaa gctggggatg aaattgtgag caatgccatc 1320
tcctacgctc tatcaaagc cttcagcacc agtgagcaag acaaggataa ctggaatggg 1380
cagctgaagc ttctgctgga gtggaaccag ctggacttag ccaatgatga gattttcacc 1440
aatgaccgcc gatgggagtc tgctgacctt caagaagtca tgtttacggc tctcataaag 1500
gacagaccca agtttgtccg cctctttctg gagaatggct tgaacctacg gaagtttctc 1560
acccatgatg tcctcactga actcttctcc aaccttca gcacgcttgt gtaccggaat 1620
ctgcagatcg ccaagaattc ctataatgat gccctctca cgtttgtctg gaaactgggt 1680
gcgaacttcc gaagaggctt ccggaaggaa gacagaaatg gccgggacga gatggacata 1740
gaactccacg acgtgtctcc tattactcgg caccctctgc aagctctct catctggggc 1800
attcttcaga ataagaagga actctccaaa gtcatttggg agcagaccag gggctgcact 1860
ctggcagccc ttggagccag caagcttctg aagactctgg ccaaagtga gaacgacatc 1920
aatgctgctg gggagtccga ggagctggct aatgagtac agaccgggc tgttgagctg 1980
ttcactgagt gttacagcag cgatgaagac ttggcagaac agctgctggg ctattcctgt 2040
gaagcttggg gtggaagcaa ctgtctggag ctggcggtgg agggcacaga ccagcatttc 2100
atgcgccagc ctggggtcca gaattttctt tctaagcaat ggtatggaga gatttcccga 2160
gacaccaaga actggaagat tatectgtgt ctgtttatta tacccttggg gggctgtggc 2220
tttgtatcat ttaggaagaa acctgtcgac aagcacaaga agctgctttg gtactatgtg 2280
gcgttcttca cctccccctt cgtggctctt tectggaatg tggctcttca catcgcttc 2340
ctcctgctgt tgcctacgt gctgtcatg gatttccatt cggtgccaca cccccctag 2400
ctggctcctg actcgtggg ctttgtcttc ttctgtgatg aagtgaagca gtggtagta 2460
aatgggggtg attattttac tgacctgtgg aatgtgatgg acacgctggg gcttttttac 2520
ttcatagcag gaattgtatt tcggctccac tcttctaata aaagctcttt gtattctgga 2580
cgagtcatct tctgtctgga ctacattatt ttcactctaa gattgatcca cattttact 2640
gtaagcagaa acttaggacc caagattata atgctgcaga ggatgctgat cgatgtgttc 2700
ttcttctgt tctctttgc ggwtggatg gtggcctttg gcgtggccag gcaagggatc 2760
cttaggcaga atgagcagcg ctggagggtg atattccgtt cggctcatct cagccctac 2820
ctggccatgt tcggccaggt gccagtgac gtggatggt ccacgtatga ctttggccac 2880
tgcaacttca ctgggaatga gtccaagcca ctgtgtgtgg agctggatga gcacaacctg 2940
ccccggttcc ccgagtggat caccatcccc ctggtgtgca tctacatgtt atccaccaac 3000
atcctgctgg tcaacctgct ggtcgccatg tttggctaca cgggtgggcac cgtccaggag 3060
aacaatgacc aggtctggaa gttccagagg tacttctgg tgcaggagta ctgagccgc 3120
ctcaatatcc cctccccctt catcgtcttc gcttacttct acatgggtgg gaagaagtgc 3180
ttcaagtgtt gctgcaagga gaaaaacatg gagtcttctg tctgctgttt caaaaatgaa 3240
gacaatgaga ctctggcatg ggagggtgtc atgaaggaaa actacctgt caagatcaac 3300
acaaaagcca acgacacctc agaggaaatg aggcacgat ttagacaact ggatacaaat 3360
cttaatgatc tcaagggtct tctgaaagag attgtctaata aaatcaata aactgtatg 3420
aactctaatt gagaaaaatc taattatagc aagatcatat taaggaaatgc tgatgaacaa 3480
ttttgctatc gactactaaa tgagagattt tcagaccctt ggggtacatg tggatgattt 3540
taaatcacc tagtgtgtg agaccttgag aataaagtgt gaaggcgaa ttctgcagat 3600
atccatcaca ctggcgccg ctcgagcatg catctagag 3639

```

<210> 8

<211> 3101

<212> DNA

<213> Homo sapiens

<400> 8

```

atggttggag gatgcagggtg gacagaagac gtggagcctg cagaagtaaa ggaaaagatg 60
tcctttcggg cagccagggt cagcatgagg aacagaagga atgacactct ggacagcacc 120
cggaccctgt actccagcgc gtctcggagc acagacttgt cttacagtga aagcgccagc 180
ttctacgtg ccttcaggac acagacgtgc ccaatcatgg cttcttggga cttggtgaat 240
tttattcaag caaattttaa gaaacgagaa tgtgtcttct ttaccaaaaga ttccaaggcc 300
acggagaatg tgtgcaagtg tggctatgcc cagagccagc acatggaagg caccagatc 360
aaccaaagtg agaaatggaa ctacaagaaa cacaccaagg aatttcctac cgacgccttt 420
ggggatattc agtttgagac actggggaag aaagggaagt atatacgtct gtcctgcgac 480
acggacgcgg aaatccttta cgagctgctg acccagcact ggcacctgaa aacaccaaac 540
ctggtcattt ctgtgaccgg gggcgccaag aacttcgccc tgaagccgag catgcgcaag 600
atcttcagcc ggctcatcta catcgcgag tccaaagggt cttggattct cacgggaggc 660
acccattatg gcctgatgaa gtacatcggg gaggtggtga gagataacac catcagcagg 720
agttcagagg agaattttgt ggccattggc atagcagctt ggggcatggt ctccaaccgg 780
gacaccctca tcaggaattg cgatgctgag ggctattttt tagcccagta ccttatggat 840
gacttcacaa gagatccact gtatatcctg gacaacaacc acacacattt gctgctcgtg 900
gacaatggct gtcattggaa tcccactgtc gaagcaaaag tccggaatca gctagagaag 960
tatatctctg agcgcactat tcaagattcc aactatgggt gcaagatccc cattgtgtgt 1020
tttgcccaag gaggtggaag agagactttg aaagccatca atacctccat caaaaataaa 1080
attccttgtg tgggtggtgga aggtcggggc cagatcgctg agtgatcgc tagcctgggt 1140
gaggtggagg atgccctgac atcttctgcc gtcaaggaga agctggtgag ctttttacc 1200
cgcacgggtg ccggctgccc tgaggaggag attagagatt ggatcaaagt gctcaaagaa 1260
attctcgaat gttctcacct attaacagtt attaaaatgg aagaagctgg ggatgaaatt 1320
gtgagcaatg ccatctccta cgctctatac aaagccttca gcaccagtga gcaagacaag 1380
gataactgga atgggcagct gaagcttctg ctggagtgga accagctgga cttagccaat 1440
gatgagattg tcaccaatga ccgcatgag gagaagagca aaccgaggct cagagacaca 1500
ataatccagg tcacatggct ggaaaatggg agaatcaagg ttgagagcaa agatgtgact 1560
gacggcaaaag cctcttctca tatgctggtg gttctcaagt ctgctgacct tcaagaagtc 1620
atgtttacgg ctctcataaa ggacagaccc aagtttgtcc gcctctttct ggagaatggc 1680
ttgaacctat ggaagtttct caccatgat gtcctcactg aactcttctc caaccacttc 1740
agcagccttg tgtaccgga tctgcagatc gccaaagaatt cctataatga tgccctcctc 1800
acgtttgtct ggaaactggg tgcgaacttc cgaagaggct tccggaagga agacagaaat 1860
ggccgggagc agatggacat agaactccac gacgtgtctc ctattactcg gcacccctg 1920
caagctctct tcactctggc cattcttcag aataagaagg aactctccaa agtcatttgg 1980
gagcagacca ggggctgccc tctggcagcc ctgggagcca gcaagcttct gaagactctg 2040
gccaaagtga agaacgacat caatgctgct ggggagtcgg aggagctggc taatgagtac 2100
gagaccgggg ctgttggtga gtccacagtg tggaaatgctg tgggtgggagc ggatctgcca 2160
tgtggcacag acattgccag cggcactcat agaccagatg gtggagagct gttcactgag 2220
tgttacagca gcgatgaaga cttggcagaa cagctgctgg tctattcctg tgaagcttgg 2280
gggtggaagca actgtctgga gctggcgggt tcttaagcaa tgggtatggg agatttccc 2340
cctgggggtc agaattttct ttctaagcaa tgggtatggg agatttccc 2400
aactggaaga ttatcctgtg tctgtttatt atacccttgg tgggtgttgg ctttgtatca 2460
tttaggaaga aaacctgtcg caagcacaag aagctgcttt ggtactatgt ggcgttcttc 2520
acctccccct tctgtgtctt ctcttggaa gtgtgtctct acatcgctt cctcctgctg 2580
tttgccctacg tgctgctcat ggatttccat tgggtgccac acccccccca gctggtcctg 2640
tactcgctgg tctttgtctt cttctgtgat gaagttagac agggccggcc ggctgctccc 2700
agtgcggggc ccgccaagcc cagccaccac tctggcccg cctgcaagc aagctccaca 2760
cgcagccccg gttcccgctc acgcccactc ttccacactt ccctgcaagc tgaggggtgc 2820
agctctggcc ttggccagcc cagaaagggg ctcccacagt gcagcggtgg gctgaagggg 2880
ctcctcaagt gcggccaaag tgggagccca ggcagaggag gcgcccagag tgagcgaggg 2940
ctgtgaggac tgccagcaag cacgtgtca cctctcagaa gcgtaagaca gcaatggacc 3000
aaacagacga agatctcttc ccctatggag cattctacca gttcctgatg atttccagga 3060
gctttcgagg agaggagatg agcatcggca agcagcacta a 3101

```

<210> 9

<211> 930

<212> DNA

<213> Homo sapiens

<400> 9

```

ggcacgaggc tgcctttctc caccagagac tcttctcag ggaggacttg gtgaatttta 60
ttcaagcaaa ttttaagaaa cgagaatgtg tcttctttac aagggcacgc 120
tcaatgaaat ccttcttccc tgtccacacc atcgtgctta tcagggagaa tgtgtgcaag 180
tgtggctatg ccagagacca gcacatggaa ggcacccaga tcaaccaaag tgagaaatgg 240
aactacaaga aacacaccaa ggaatttctt accgacgcct ttggggatat tcagttttag 300
acactgggga agaaagggaa gtatatagct ctgtcctgag acacggacgc ggaatcctt 360

```

| | | | | | | |
|------------|-------------|------------|------------|-------------|------------|-----|
| tacgagctgc | tgacccagca | ctggcacctg | aaaacaccca | acctgggtcat | ttctgtgacc | 420 |
| gggggcgcca | agaacttcgc | cctgaagccg | cgcatgcgca | agatcttcag | ccggctcatc | 480 |
| tacatcgcg | agtccaaagg | tgcttggtat | ctcacgggag | gcacccatta | tggcctgatg | 540 |
| aagtacatcg | gggaggtggt | gagagataac | accatcagca | ggagttcaga | ggagaatatt | 600 |
| gtggccattg | gcatagcagc | ttggggcatg | gtctccaacc | gggacaccct | catcaggaat | 660 |
| tgcgatgctg | aggtagccgt | gggacaggag | gaggtctgct | aggtcacatg | gaagaaagac | 720 |
| catggcatgg | gcctgtggcc | tgaaccctgg | ggctctgtga | tggagccagc | cagatcatgg | 780 |
| ggaagtctgc | ctttcaagga | gtgcctttgg | gaccttaaag | gaattgaaaa | caaggatgac | 840 |
| gtacctaat | aactgctggg | aaagagttaa | caatgaatgt | tttgttcatt | aaaatgtgtt | 900 |
| ctcagcaatc | tcaaaaaaaaa | aaaaaaaaaa | | | | 930 |

<210> 10

<211> 3288

<212> DNA

<213> Homo sapiens

<400> 10

| | | | | | | |
|-------------|-------------|-------------|-------------|-------------|-------------|------|
| atgaggaaca | gaaggaatga | cactctggac | agcaccggga | ccctgtactc | cagcgcgtct | 60 |
| cggagcagag | acttgtctta | cagtgaagac | gacttggtga | atattattca | agcaaatttt | 120 |
| aagaaacgag | aatgtgtctt | ctttaccaa | gattccaagg | ccacggagaa | tgtgtgcaag | 180 |
| tgtggctatg | cccagagcca | gcacatggaa | ggcaccgaga | tcaaccaaag | tgagaaatgg | 240 |
| aactacaaga | aacacaccaa | ggaatttcct | accgacgcct | ttggggatat | tcagtttgag | 300 |
| acactgggga | agaaagggaa | gtatatacgt | ctgtcctgcg | acacggacgc | ggaaatcctt | 360 |
| tacgagctgc | tgacccagca | ctggcacctg | aaaacaccca | acctgggtcat | ttctgtgacc | 420 |
| gggggcgcca | agaacttcgc | cctgaagccg | cgcatgcgca | agatcttcag | ccggctcatc | 480 |
| tacatcgcg | agtccaaagg | tgcttggtat | ctcacgggag | gcacccatta | tggcctgatg | 540 |
| aagtacatcg | gggaggtggt | gagagataac | accatcagca | ggagttcaga | ggagaatatt | 600 |
| gtggccattg | gcatagcagc | ttggggcatg | gtctccaacc | gggacaccct | catcaggaat | 660 |
| tgcgatgctg | agggtctattt | tttagccag | taccttatgg | atgacttcac | aagagatcca | 720 |
| ctctatatcc | tggacaacaa | ccacacacat | ttgctgctcg | tggacaatgg | ctgtcatgga | 780 |
| catcccactg | tcgaagcaaa | gctccggaat | cagctagaga | agtatatctc | tgagcgcact | 840 |
| attcaagatt | ccaactatgg | tggcaagatc | cccattgtgt | gttttgccca | aggaggtgga | 900 |
| aaagagactt | tgaaagccat | caatacctcc | atcaaaaaata | aaattccttg | tgtggtgggtg | 960 |
| gaaggctcgg | gccagatcgc | tgatgtgatc | gctagcctgg | tggaggtgga | ggatgccctg | 1020 |
| acatcttctg | cgttcaagga | gaagctgggtg | cgctttttac | cccgcacggg | gtcccggctg | 1080 |
| cctgaggagg | agactgagag | ttggatcaaa | tggctcaaag | aaattctcga | atgttctcac | 1140 |
| ctattaacag | ttattaaaat | ggaagaagct | ggggatgaaa | ttgtgagcaa | tgccatctcc | 1200 |
| tacgtcttat | acaaagcctt | cagcaccagt | gagcaagaca | aggataactg | gaatgggcag | 1260 |
| ctgaagcttc | tgctggagtg | gaaccagctg | gacttagcca | atgatgagat | tttcaccagac | 1320 |
| gaccgccgat | gggagtctgc | tgaccttcaa | gaagtcatgt | ttacggctct | cataaaggac | 1380 |
| agacccaagt | ttgtccgcct | ctttctggag | aatggcttga | acctacggaa | gtttctcacc | 1440 |
| catgatgtcc | tacttgaact | cttctccaac | cacttcagca | cgcttgtgta | ccggaatctg | 1500 |
| cagatcgcca | agaattccta | taatgatgcc | ctcctcacgt | ttgtctggaa | actggttgcg | 1560 |
| aaacttcgaa | gaggcttccg | gaaggaagac | agaaatggcc | gggacgagat | ggacatagaa | 1620 |
| ctccacgacg | tgtctcctat | tactcggcac | cccctgcaag | ctctcttcat | ctgggccatt | 1680 |
| cttcagaata | agaaggaact | ctccaaagtc | atltgggagc | agaccagggg | ctgcaactctg | 1740 |
| gcagccctgg | gagccagcaa | gcttctgaag | actctggcca | aagtgaagaa | cgacatcaat | 1800 |
| gctgttgggg | agtcggagga | gctggcta | gagtaacgga | cccgggctgt | tgagctgttc | 1860 |
| actgagtgtt | acagcagcga | tgaagacttg | gcagaacagc | tgctggtcta | ttcctgtgaa | 1920 |
| gcttgggggtg | gaagcaactg | tctggagctg | gcgggtggag | ccacagacca | gcatttcatc | 1980 |
| gcccagcctg | gggtccagaa | ttttctttct | aagcaatggt | atggagagat | ttcccagac | 2040 |
| accaagaacc | ggaagattat | cctgtgtctg | tttattatac | ccttgggtgg | ctgtggcttt | 2100 |
| gtatcattta | ggaagaaacc | tgtcgacaag | cacaagaagc | tgctttggta | ctatgtggcg | 2160 |
| ttcttcacct | cccccttctg | ggtcttctcc | tggaaatgtg | tcttctacat | cgcttctctc | 2220 |
| ctgctgtttg | cctacgtgct | gctcatggat | ttccattcgg | tgccacaccc | ccccagctg | 2280 |
| gtcctgtact | cgtgtgtctt | tgtcctcttc | tgtcatgaag | tgagacagtg | tcacgttaatt | 2340 |
| gggggtgaatt | atlttactga | cctgtggaat | gtgatggaca | cgctggggct | tttttacttc | 2400 |
| atagcaggaa | ttgtatttctg | gctccactct | tctaataaaa | gctctttgta | ttctggacga | 2460 |
| gtcattttct | gtctggacta | cattattttc | actctaagat | tgatccacat | ttttactgta | 2520 |
| agcagaaacc | taggacccaa | gattataatg | ctgcagagga | tgctgatcga | tgtgttcttc | 2580 |
| ttcctgttcc | tctttgcggg | gtggatgggtg | gcctttggcg | tggccaggca | agggatcctt | 2640 |
| aggcagaatg | agcagcgtg | gaggtggata | ttccgttcgg | tcactctacga | gccctacctg | 2700 |
| gccatgttctg | gccaggtgcc | cagtgcagtg | gatggtacca | cgtatgactt | tgcccactgc | 2760 |
| accttccactg | ggaatgagtc | caagccactg | tgtgtggagc | tggatgagca | caacctgcc | 2820 |
| cggttccccg | agtggatcac | catccccctg | gtgtgcatct | acatgttatc | caccaacatc | 2880 |
| ctgctggtca | acctgctggt | cgccatgttt | ggctacacgg | tgggcaccgt | ccaggagAAC | 2940 |
| aatgaccagg | tctggaagtt | ccagaggtac | ttcctggtgc | aggagtactg | cagccgcctc | 3000 |

- 11 -

```

aatatcccct tccccttcat cgtcttcgct tacttctaca tgggtggtgaa gaagtgtctc 3060
aagtgttgct gcaaggagaa aaacatggag tcttctgtct gctgtttcaa aaatgaagac 3120
aatgagactc tggcatggga ggggtgtcatg aaggaaaact accttgtcaa gatcaacaca 3180
aaagccaacg acacctcaga ggaaatgagg catcgattta gacaactgga tacaagactt 3240
aatgatctca agggctctct gaaagagatt gctaataaaa tcaataaa 3288

```

<210> 11

<211> 5674

<212> DNA

<213> Homo sapiens

<400> 11

```

aagaaaaatcc tgcttgacaa aaaccgtcac ttaggaaaag atgtcctttc gggcagccag 60
gctcagcatg aggaacagaa ggaatgacac tctggacagc acccggaccc tgtactccag 120
cgcgctctcg agcacagact tgtcttacag tgaaagcgac ttggtgaatt ttattcaagc 180
aaattttaag aaacgagaat gtgtcttctt tatcaaagat tccaaggcca cggagaatgt 240
gtgcaagtgt ggctatgccc agagccagca catggaaggc acccagatca accaaagtga 300
gaaatggaac tacaagaaac acaccaagga atttcctacc gacgcctttg gggatattca 360
gtttgagaca ctggggaaga aagggaagta tatacgtctg tctgtcgaca cggacgcgga 420
aatcctttac gagctgctga cccagcactg gcacctgaaa acacccacc tggatatttc 480
tgtgaccggg ggcgccaaga acttcgccct gaagccgcgc atgcgcaaga tcttcagccg 540
gctcatctac atcgcgagc ccaaaggctg ttggattctc acgggaggca cccattatgg 600
cctgatgaag tacatcgggg aggtggtgag agataacacc atcagcagga gttagagga 660
gaatattgtg gccattggca tagcagcttg gggcatggtc tccaaccggg acacctcat 720
caggaattgc gatgctgagg gctatttttt agcccagtag cttatggatg acttcacaag 780
agatccactg tatatcctgg acaacaacca cacacatttg ctgctcgtag acaatggctg 840
tcatggacat cccactgtcg aagcaaagct ccggaatcag cttagagaag atattctctga 900
gcgcactatt caagattcca actatggtgg caagatcccc attgtgtgtt ttgcccagg 960
aggtggaaaa gagactttga aagccatcaa tactctcatc aaaaataaaa ttccttgtgt 1020
ggtggtggaa ggctcgggcc agatcgctga tctgatcgct agcctggtgg aggtggaggga 1080
tgccctgaca tcttctgccc tcaaggagaa gctggtgcgc tttttacccc gcacggtgtc 1140
ccggctgcct gaggaggaga ctgagagttg gatcaaattg ctcaaagaaa ttctcgaatg 1200
ttctcaccta ttaacagtta ttaaaatgga agaagctggg gatgaaattg tgagcaatgc 1260
catctcctac atgctctaca aagccttcag ccagctggag caagacaagg ataactggaa 1320
tgggcagctg aagcttctgc tggagtggaa ccagctggac ttagccaatg atgagatttt 1380
caccaatgac cgccgatggg agtctgctga ccttcaagaa gtcatgttta cggctctcat 1440
aaaggacaga cccaagtttg tccgcctctt tctggagaat ggctgaacc tacggaagtt 1500
tctcaccat atgtctctca ctgaactctt ctccaaccac ttcagcacgc ttgtgtaccg 1560
gaatctgcag atcgccaaga attcctataa tgatgccctc ctcacgtttg tctggaaact 1620
ggttgcgaa ttccgaagag gcttccggaa ggaagacaga aatggccggg acgagatgga 1680
catagaactc cagcagctgt ctctatttac tgggaccccc ctgcaagctc tcttcatctg 1740
ggccattctt cagaataaga aggaactctc caaagtcatt tgggagcaga ccaggggctg 1800
cactctggca gccctgggag ccagcaagct tctgaagact ctggccaaag tgaagaacga 1860
catcaatgct gctggggagt ccgaggagct ggctaagtag tacgagaccc gggctgttga 1920
gctgttcaact gagtgttaca gcagcgatga agacttggca gaacagctgc tggcttattc 1980
ctgtgaagct tggggtggaa gcaactgtct ggaagtgccg gtggaggcca cagaccagca 2040
tttcatcgcc cagcctgggg tccagaattt tctttctaag caatggtagt gagagatttc 2100
ccgagacacc aagaactgga agattatcct gtgtctgttt attataccct tgggtgggctg 2160
tggctttgta tcatttagga agaaacctgt cgacaagcac aagaagctgc tttggtacta 2220
tgtggcgctt ttcacctccc ccttcgtggt ctctcctcgg aatgtggtct tctacatcgc 2280
cttccctcct ctggttgccct acgtgctgct catggatttc cattcggtgc cacaccccc 2340
cgagctggtc ctgtactcgc tggctcttgt cctcttctgt gatgaagtga gacagtggta 2400
cgtaaatggg gtgaattatt ttactgacct gtggaatgtg atggacacgc tggggctttt 2460
ttacttcata gcaggaattg tatttcggct tggactacat tattttcact ctaagattga tccacatttt 2580
tggacagatc agaaacttag gacccaagat tataatgctg cagaggatgc tgatcgatgt 2640
gttcttcttc ctgttccctt ttgctgtgtg gatgggtggc tttggcggtg ccaggcaagg 2700
gatccttagg cagaatgagc agcgtctggg gtggaatttc cgttcgggca tctacgagcc 2760
ctacctggcc atgttcggcc aggtgccag ttccagattt cgttaccacgt atgactttgc 2820
ccactgcacc ttactggga atgagtccaa gccactgtgt gtggagctgg atgagcaca 2880
cctgccccgg ttccccgagt ggatcaccat cccctggtg tgcattctaca tgttatccac 2940
caacatccct ctggtcaacc tgctggtcgc catgtttggc tacacggtgg gcaccgtcca 3000
ggagaaccaat gaccaggtct ggaagttcca gaggtaactc ctgggtgcagg agtactgcag 3060
ccgcctcaat atccccctcc ccttcatcgt cttcgcttac ttctacatgg tgggaagaa 3120
gtgcttcaag tgttgctgca aggagaaaaa catggagtct tctgtctgct gtttcaaaaa 3180
tgagacaat gagactctgg catgggaggg tgtcatgaag gaaaactacc ttgtcaagat 3240
caacacaaaa gccaacgaca cctcagagga aatgaggcat cgatttagac aactggatac 3300

```

- 12 -

```

aaagcttaaat gatctcaagg gtcttctgaa agagattgct aataaaatca aataaaatca 3360
aataaaactg tatgaaactc taatggagaa aaatctaatt atagcaagat catattaagg 3420
aatgctgatg aacaattttg ctatcgacta ctaaatagaga gattttcaga cccctgggta 3480
catgggtggat gatttttaaat caccctagtg tgctgagaoc ttgagaataa agtgtgtgat 3540
tggtttcata cttgaagacg gatataaagg aagaataatt cctttatgtg tttctccaga 3600
atgggtgctg tttctctctg tgtctcaatg cctgggactg gaggttgata gtttaagtgt 3660
gttctttaccg cctccttttt cctttaatct tatttttgat gaacacatat ataggagaac 3720
atctatccta tgaataagaa cctgggtcatg ctttactcct gtattgttat tttgttcatt 3780
tccaattgat tctctacttt tccctttttt gtattatgtg actaattagt tggcatattg 3840
ttaaaagtct ctcaaattag gccagattct aaaaactgct gcagcaagag gaccccgctc 3900
tcttcaggaa aagtgttttc atttctcagg atgcttctta cctgtcagag gaggtgacaa 3960
ggcagtgctct tgctctcttg gactcaccag gctcctattg aaggaaccac cccctattcct 4020
aaatatgtga aaagtcgccc aaaatgcaac cttgaaaggc actactgact ttgttcttat 4080
tggatactcc tcttattttat tatttttcca ttaaaaataa tagctggcta ttatagaaaa 4140
tttagaccat acagagatgt agaaagaaca taaattgtcc ccattacctt aagggtatca 4200
ctgctaacaa tttctggatg gtttttcaag tctatttttt ttctatgtat gtctcaattc 4260
tctttcaaaa ttttacagaa tgttatcata ctacatatat acttttttat taagcttttt 4320
cacttagtat tttatcaaat atgtttttat tatattcata gccttcttaa acattatatc 4380
aataattgca taataggcaa cctctagcga ttaccataat tttgctcatt gaaggctatc 4440
tccagttgat cattgggatg agcatctttg tgcataaacc ctattgctgt atttgggaaa 4500
atfttccaag gttagattcc aataaatatc tatttattat taaatattaa aatatcgatt 4560
tattattaaa accattttata aggtcttttc ataatgtat agcaaatagg aattattaac 4620
ttgagcataa gatagagat acatgaacct gaactattaa aataaaatat tatatttaac 4680
cctagtttaa gaagaagtca atatgcttat ttaaatatta tggatgggtg gcagatcact 4740
tgaggtcagg agttcgagac cagcctggcc aacatggcaa aaccacatct ctactaaaaa 4800
taaaaaaatt agctgggtgt ggtgggtgcac tctgtaatc ccagctactc agaaggctga 4860
ggtacaagaa ttgctggaac ctgggagggc gaggttgag tgaaccaaga ttgcaccact 4920
gcactccagc cggggtgaca gactgagact ccgactgaaa ataaataaat aaataaataa 4980
ataaataaat aaataaatat tatggatggt cccagctttc totggaagtg gtggtatttg agcaggatgt 5100
tcttactgaa cacctgtagt ccataattag tttctcagct ttgaatacac tataaactca 5160
gcacaaggca attgaaatgc tagaaggaa ctactaaaag atctaatttg aaaaactaca 5220
gtggctgaag gaggaatttt tagaaggaa ctactaaaag atctaatttg aaaaactaca 5220
aaagcattaa ctaaaaaagt ttattttcct tttgtctggg cagtagtgaa aataactact 5280
cacaacattc actatgtttg caaggaaatc acacaaataa aagatgcctt tttacttaaa 5340
cgccaagaca gaaaacttgc ccaatactga gaagcaactt gcattagaga gggaaactgt 5400
aaatgttttc aaccagttc atctggtgga tgtttttgca ggttactctg agaattttgc 5460
ttatgaaaaa tcattatttt tagtgtagtt cacaataatg tattgaacat acttctaatt 5520
aaaggtgcta tgccttctgt tatggtacta aatgtgtcct gtgtactttt gcacaactga 5580
gaatcctgag gcttggttta atgagtgtgt tcatgaaata aataatggag gaattgtcaa 5640
aaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaa 5674

```

<210> 12
 <211> 1095
 <212> PRT
 <213> Homo sapiens

<400> 12
 Met Arg Asn Arg Arg Asn Asp Thr Leu Asp Ser Thr Arg Thr Leu Tyr
 1 5 10 15
 Ser Ser Ala Ser Arg Ser Thr Asp Leu Ser Tyr Ser Glu Ser Asp Leu
 20 25 30
 Val Asn Phe Ile Gln Ala Asn Phe Lys Lys Arg Glu Cys Val Phe Phe
 35 40 45
 Thr Lys Asp Ser Lys Ala Thr Glu Asn Val Cys Lys Cys Gly Tyr Ala
 50 55 60
 Gln Ser Gln His Met Glu Gly Thr Gln Ile Asn Gln Ser Glu Lys Trp
 65 70 75 80
 Asn Tyr Lys Lys His Thr Lys Glu Phe Pro Thr Asp Ala Phe Gly Asp
 85 90 95
 Ile Gln Phe Glu Thr Leu Gly Lys Lys Gly Lys Tyr Ile Arg Leu Ser
 100 105 110

Cys Asp Thr Asp Ala Glu Ile Leu Tyr Glu Leu Leu Thr Gln His Trp
 115 120 125
 His Leu Lys Thr Pro Asn Leu Val Ile Ser Val Thr Gly Gly Ala Lys
 130 135 140
 Asn Phe Ala Leu Lys Pro Arg Met Arg Lys Ile Phe Ser Arg Leu Ile
 145 150 155 160
 Tyr Ile Ala Gln Ser Lys Gly Ala Trp Ile Leu Thr Gly Gly Thr His
 165 170 175
 Tyr Gly Leu Met Lys Tyr Ile Gly Glu Val Val Arg Asp Asn Thr Ile
 180 185 190
 Ser Arg Ser Ser Glu Glu Asn Ile Val Ala Ile Gly Ile Ala Ala Trp
 195 200 205
 Gly Met Val Ser Asn Arg Asp Thr Leu Ile Arg Asn Cys Asp Ala Glu
 210 215 220
 Gly Tyr Phe Leu Ala Gln Tyr Leu Met Asp Asp Phe Thr Arg Asp Pro
 225 230 235 240
 Leu Tyr Ile Leu Asp Asn Asn His Thr His Leu Leu Leu Val Asp Asn
 245 250 255
 Gly Cys His Gly His Pro Thr Val Glu Ala Lys Leu Arg Asn Gln Leu
 260 265 270
 Glu Lys Tyr Ile Ser Glu Arg Thr Ile Gln Asp Ser Asn Tyr Gly Gly
 275 280 285
 Lys Ile Pro Ile Val Cys Phe Ala Gln Gly Gly Gly Lys Glu Thr Leu
 290 295 300
 Lys Ala Ile Asn Thr Ser Ile Lys Asn Lys Ile Pro Cys Val Val Val
 305 310 315 320
 Glu Gly Ser Gly Gln Ile Ala Asp Val Ile Ala Ser Leu Val Glu Val
 325 330 335
 Glu Asp Ala Leu Thr Ser Ser Ala Val Lys Glu Lys Leu Val Arg Phe
 340 345 350
 Leu Pro Arg Thr Val Ser Arg Leu Pro Glu Glu Glu Thr Glu Ser Trp
 355 360 365
 Ile Lys Trp Leu Lys Glu Ile Leu Glu Cys Ser His Leu Leu Thr Val
 370 375 380
 Ile Lys Met Glu Glu Ala Gly Asp Glu Ile Val Ser Asn Ala Ile Ser
 385 390 395 400
 Tyr Ala Leu Tyr Lys Ala Phe Ser Thr Ser Glu Gln Asp Lys Asp Asn
 405 410 415
 Trp Asn Gly Gln Leu Lys Leu Leu Leu Glu Trp Asn Gln Leu Asp Leu
 420 425 430
 Ala Asn Asp Glu Ile Phe Thr Asn Asp Arg Arg Trp Glu Ser Ala Asp
 435 440 445
 Leu Gln Glu Val Met Phe Thr Ala Leu Ile Lys Asp Arg Pro Lys Phe
 450 455 460

- 14 -

Val Arg Leu Phe Leu Glu Asn Gly Leu Asn Leu Arg Lys Phe Leu Thr
 465 470 475 480
 His Asp Val Leu Thr Glu Leu Phe Ser Asn His Phe Ser Thr Leu Val
 485 490 495
 Tyr Arg Asn Leu Gln Ile Ala Lys Asn Ser Tyr Asn Asp Ala Leu Leu
 500 505 510
 Thr Phe Val Trp Lys Leu Val Ala Asn Phe Arg Arg Gly Phe Arg Lys
 515 520 525
 Glu Asp Arg Asn Gly Arg Asp Glu Met Asp Ile Glu Leu His Asp Val
 530 535 540
 Ser Pro Ile Thr Arg His Pro Leu Gln Ala Leu Phe Ile Trp Ala Ile
 545 550 555 560
 Leu Gln Asn Lys Lys Glu Leu Ser Lys Val Ile Trp Glu Gln Thr Arg
 565 570 575
 Gly Cys Thr Leu Ala Ala Leu Gly Ala Ser Lys Leu Leu Lys Thr Leu
 580 585 590
 Ala Lys Val Lys Asn Asp Ile Asn Ala Ala Gly Glu Ser Glu Glu Leu
 595 600 605
 Ala Asn Glu Tyr Glu Thr Arg Ala Val Glu Leu Phe Thr Glu Cys Tyr
 610 615 620
 Ser Ser Asp Glu Asp Leu Ala Glu Gln Leu Leu Val Tyr Ser Cys Glu
 625 630 635 640
 Ala Trp Gly Gly Ser Asn Cys Leu Glu Leu Ala Val Glu Ala Thr Asp
 645 650 655
 Gln His Phe Ile Ala Gln Pro Gly Val Gln Asn Phe Leu Ser Lys Gln
 660 665 670
 Trp Tyr Gly Glu Ile Ser Arg Asp Thr Lys Asn Arg Lys Ile Ile Leu
 675 680 685
 Cys Leu Phe Ile Ile Pro Leu Val Gly Cys Gly Phe Val Ser Phe Arg
 690 695 700
 Lys Lys Pro Val Asp Lys His Lys Lys Leu Leu Trp Tyr Tyr Val Ala
 705 710 715 720
 Phe Phe Thr Ser Pro Phe Val Val Phe Ser Trp Asn Val Val Phe Tyr
 725 730 735
 Ile Ala Phe Leu Leu Leu Phe Ala Tyr Val Leu Leu Met Asp Phe His
 740 745 750
 Ser Val Pro His Pro Pro Glu Leu Val Leu Tyr Ser Leu Val Phe Val
 755 760 765
 Leu Phe Cys Asp Glu Val Arg Gln Trp Tyr Val Asn Gly Val Asn Tyr
 770 775 780
 Phe Thr Asp Leu Trp Asn Val Met Asp Thr Leu Gly Leu Phe Tyr Phe
 785 790 795 800
 Ile Ala Gly Ile Val Phe Arg Leu His Ser Ser Asn Lys Ser Ser Leu
 805 810 815
 Tyr Ser Gly Arg Val Ile Phe Cys Leu Asp Tyr Ile Ile Phe Thr Leu
 820 825 830

- 15 -

Arg Leu Ile His Ile Phe Thr Val Ser Arg Asn Leu Gly Pro Lys Ile
 835 840 845
 Ile Met Leu Gln Arg Met Leu Ile Asp Val Phe Phe Phe Leu Phe Leu
 850 855 860
 Phe Ala Val Trp Met Val Ala Phe Gly Val Ala Arg Gln Gly Ile Leu
 865 870 875 880
 Arg Gln Asn Glu Gln Arg Trp Arg Trp Ile Phe Arg Ser Val Ile Tyr
 885 890 895
 Glu Pro Tyr Leu Ala Met Phe Gly Gln Val Pro Ser Asp Val Asp Gly
 900 905 910
 Thr Thr Tyr Asp Phe Ala His Cys Thr Phe Thr Gly Asn Glu Ser Lys
 915 920 925
 Pro Leu Cys Val Glu Leu Asp Glu His Asn Leu Pro Arg Phe Pro Glu
 930 935 940
 Trp Ile Thr Ile Pro Leu Val Cys Ile Tyr Met Leu Ser Thr Asn Ile
 945 950 955 960
 Leu Leu Val Asn Leu Leu Val Ala Met Phe Gly Tyr Thr Val Gly Thr
 965 970 975
 Val Gln Glu Asn Asn Asp Gln Val Trp Lys Phe Gln Arg Tyr Phe Leu
 980 985 990
 Val Gln Glu Tyr Cys Ser Arg Leu Asn Ile Pro Phe Pro Phe Ile Val
 995 1000 1005
 Phe Ala Tyr Phe Tyr Met Val Val Lys Lys Cys Phe Lys Cys Cys Cys
 1010 1015 1020
 Lys Glu Lys Asn Met Glu Ser Ser Val Cys Cys Phe Lys Asn Glu Asp
 1025 1030 1035 1040
 Asn Glu Thr Leu Ala Trp Glu Gly Val Met Lys Glu Asn Tyr Leu Val
 1045 1050 1055
 Lys Ile Asn Thr Lys Ala Asn Asp Thr Ser Glu Glu Met Arg His Arg
 1060 1065 1070
 Phe Arg Gln Leu Asp Thr Lys Leu Asn Asp Leu Lys Gly Leu Leu Lys
 1075 1080 1085
 Glu Ile Ala Asn Lys Ile Lys
 1090 1095

<210> 13
 <211> 1104
 <212> PRT
 <213> Homo sapiens

<400> 13
 Met Ser Phe Arg Ala Ala Arg Leu Ser Met Arg Asn Arg Arg Asn Asp
 1 5 10 15
 Thr Leu Asp Ser Thr Arg Thr Leu Tyr Ser Ser Ala Ser Arg Ser Thr
 20 25 30
 Asp Leu Ser Tyr Ser Glu Ser Asp Leu Val Asn Phe Ile Gln Ala Asn
 35 40 45

- 16 -

Phe Lys Lys Arg Glu Cys Val Phe Phe Ile Lys Asp Ser Lys Ala Thr
 50 55 60
 Glu Asn Val Cys Lys Cys Gly Tyr Ala Gln Ser Gln His Met Glu Gly
 65 70 75 80
 Thr Gln Ile Asn Gln Ser Glu Lys Trp Asn Tyr Lys Lys His Thr Lys
 85 90 95
 Glu Phe Pro Thr Asp Ala Phe Gly Asp Ile Gln Phe Glu Thr Leu Gly
 100 105 110
 Lys Lys Gly Lys Tyr Ile Arg Leu Ser Cys Asp Thr Asp Ala Glu Ile
 115 120 125
 Leu Tyr Glu Leu Leu Thr Gln His Trp His Leu Lys Thr Pro Asn Leu
 130 135 140
 Val Ile Ser Val Thr Gly Gly Ala Lys Asn Phe Ala Leu Lys Pro Arg
 145 150 155 160
 Met Arg Lys Ile Phe Ser Arg Leu Ile Tyr Ile Ala Gln Ser Lys Gly
 165 170 175
 Ala Trp Ile Leu Thr Gly Gly Thr His Tyr Gly Leu Met Lys Tyr Ile
 180 185 190
 Gly Glu Val Val Arg Asp Asn Thr Ile Ser Arg Ser Ser Glu Glu Asn
 195 200 205
 Ile Val Ala Ile Gly Ile Ala Ala Trp Gly Met Val Ser Asn Arg Asp
 210 215 220
 Thr Leu Ile Arg Asn Cys Asp Ala Glu Gly Tyr Phe Leu Ala Gln Tyr
 225 230 235 240
 Leu Met Asp Asp Phe Thr Arg Asp Pro Leu Tyr Ile Leu Asp Asn Asn
 245 250 255
 His Thr His Leu Leu Leu Val Asp Asn Gly Cys His Gly His Pro Thr
 260 265 270
 Val Glu Ala Lys Leu Arg Asn Gln Leu Glu Lys Tyr Ile Ser Glu Arg
 275 280 285
 Thr Ile Gln Asp Ser Asn Tyr Gly Gly Lys Ile Pro Ile Val Cys Phe
 290 295 300
 Ala Gln Gly Gly Gly Lys Glu Thr Leu Lys Ala Ile Asn Thr Ser Ile
 305 310 315 320
 Lys Asn Lys Ile Pro Cys Val Val Val Glu Gly Ser Gly Gln Ile Ala
 325 330 335
 Asp Val Ile Ala Ser Leu Val Glu Val Glu Asp Ala Leu Thr Ser Ser
 340 345 350
 Ala Val Lys Glu Lys Leu Val Arg Phe Leu Pro Arg Thr Val Ser Arg
 355 360 365
 Leu Pro Glu Glu Glu Thr Glu Ser Trp Ile Lys Trp Leu Lys Glu Ile
 370 375 380
 Leu Glu Cys Ser His Leu Leu Thr Val Ile Lys Met Glu Glu Ala Gly
 385 390 395 400
 Asp Glu Ile Val Ser Asn Ala Ile Ser Tyr Ala Leu Tyr Lys Ala Phe
 405 410 415

- 17 -

Ser Thr Ser Glu Gln Asp Lys Asp Asn Trp Asn Gly Gln Leu Lys Leu
 420 425 430
 Leu Leu Glu Trp Asn Gln Leu Asp Leu Ala Asn Asp Glu Ile Phe Thr
 435 440 445
 Asn Asp Arg Arg Trp Glu Ser Ala Asp Leu Gln Glu Val Met Phe Thr
 450 455 460
 Ala Leu Ile Lys Asp Arg Pro Lys Phe Val Arg Leu Phe Leu Glu Asn
 465 470 475 480
 Gly Leu Asn Leu Arg Lys Phe Leu Thr His Asp Val Leu Thr Glu Leu
 485 490 495
 Phe Ser Asn His Phe Ser Thr Leu Val Tyr Arg Asn Leu Gln Ile Ala
 500 505 510
 Lys Asn Ser Tyr Asn Asp Ala Leu Leu Thr Phe Val Trp Lys Leu Val
 515 520 525
 Ala Asn Phe Arg Arg Gly Phe Arg Lys Glu Asp Arg Asn Gly Arg Asp
 530 535 540
 Glu Met Asp Ile Glu Leu His Asp Val Ser Pro Ile Thr Arg His Pro
 545 550 555 560
 Leu Gln Ala Leu Phe Ile Trp Ala Ile Leu Gln Asn Lys Lys Glu Leu
 565 570 575
 Ser Lys Val Ile Trp Glu Gln Thr Arg Gly Cys Thr Leu Ala Ala Leu
 580 585 590
 Gly Ala Ser Lys Leu Leu Lys Thr Leu Ala Lys Val Lys Asn Asp Ile
 595 600 605
 Asn Ala Ala Gly Glu Ser Glu Glu Leu Ala Asn Glu Tyr Glu Thr Arg
 610 615 620
 Ala Val Glu Leu Phe Thr Glu Cys Tyr Ser Ser Asp Glu Asp Leu Ala
 625 630 635 640
 Glu Gln Leu Leu Val Tyr Ser Cys Glu Ala Trp Gly Gly Ser Asn Cys
 645 650 655
 Leu Glu Leu Ala Val Glu Ala Thr Asp Gln His Phe Ile Ala Gln Pro
 660 665 670
 Gly Val Gln Asn Phe Leu Ser Lys Gln Trp Tyr Gly Glu Ile Ser Arg
 675 680 685
 Asp Thr Lys Asn Trp Lys Ile Ile Leu Cys Leu Phe Ile Ile Pro Leu
 690 695 700
 Val Gly Cys Gly Phe Val Ser Phe Arg Lys Lys Pro Val Asp Lys His
 705 710 715 720
 Lys Lys Leu Leu Trp Tyr Tyr Val Ala Phe Phe Thr Ser Pro Phe Val
 725 730 735
 Val Phe Ser Trp Asn Val Val Phe Tyr Ile Ala Phe Leu Leu Leu Phe
 740 745 750
 Ala Tyr Val Leu Leu Met Asp Phe His Ser Val Pro His Pro Pro Glu
 755 760 765

- 18 -

Leu Val Leu Tyr Ser Leu Val Phe Val Leu Phe Cys Asp Glu Val Arg
 770 775 780
 Gln Trp Tyr Val Asn Gly Val Asn Tyr Phe Thr Asp Leu Trp Asn Val
 785 790 795 800
 Met Asp Thr Leu Gly Leu Phe Tyr Phe Ile Ala Gly Ile Val Phe Arg
 805 810 815
 Leu His Ser Ser Asn Lys Ser Ser Leu Tyr Ser Gly Arg Val Ile Phe
 820 825 830
 Cys Leu Asp Tyr Ile Ile Phe Thr Leu Arg Leu Ile His Ile Phe Thr
 835 840 845
 Val Ser Arg Asn Leu Gly Pro Lys Ile Ile Met Leu Gln Arg Met Leu
 850 855 860
 Ile Asp Val Phe Phe Phe Leu Phe Leu Phe Ala Val Trp Met Val Ala
 865 870 875 880
 Phe Gly Val Ala Arg Gln Gly Ile Leu Arg Gln Asn Glu Gln Arg Trp
 885 890 895
 Arg Trp Ile Phe Arg Ser Val Ile Tyr Glu Pro Tyr Leu Ala Met Phe
 900 905 910
 Gly Gln Val Pro Ser Asp Val Asp Gly Thr Thr Tyr Asp Phe Ala His
 915 920 925
 Cys Thr Phe Thr Gly Asn Glu Ser Lys Pro Leu Cys Val Glu Leu Asp
 930 935 940
 Glu His Asn Leu Pro Arg Phe Pro Glu Trp Ile Thr Ile Pro Leu Val
 945 950 955 960
 Cys Ile Tyr Met Leu Ser Thr Asn Ile Leu Leu Val Asn Leu Leu Val
 965 970 975
 Ala Met Phe Gly Tyr Thr Val Gly Thr Val Gln Glu Asn Asn Asp Gln
 980 985 990
 Val Trp Lys Phe Gln Arg Tyr Phe Leu Val Gln Glu Tyr Cys Ser Arg
 995 1000 1005
 Leu Asn Ile Pro Phe Pro Phe Ile Val Phe Ala Tyr Phe Tyr Met Val
 1010 1015 1020
 Val Lys Lys Cys Phe Lys Cys Cys Cys Lys Glu Lys Asn Met Glu Ser
 1025 1030 1035 1040
 Ser Val Cys Cys Phe Lys Asn Glu Asp Asn Glu Thr Leu Ala Trp Glu
 1045 1050 1055
 Gly Val Met Lys Glu Asn Tyr Leu Val Lys Ile Asn Thr Lys Ala Asn
 1060 1065 1070
 Asp Thr Ser Glu Glu Met Arg His Arg Phe Arg Gln Leu Asp Thr Lys
 1075 1080 1085
 Leu Asn Asp Leu Lys Gly Leu Leu Lys Glu Ile Ala Asn Lys Ile Lys
 1090 1095 1100

- 19 -

<210> 14
 <211> 192
 <212> PRT
 <213> Homo sapiens

<400> 14
 Met Lys Ser Phe Leu Pro Val His Thr Ile Val Leu Ile Arg Glu Asn
 1 5 10 15
 Val Cys Lys Cys Gly Tyr Ala Gln Ser Gln His Met Glu Gly Thr Gln
 20 25 30
 Ile Asn Gln Ser Glu Lys Trp Asn Tyr Lys Lys His Thr Lys Glu Phe
 35 40 45
 Pro Thr Asp Ala Phe Gly Asp Ile Gln Phe Glu Thr Leu Gly Lys Lys
 50 55 60
 Gly Lys Tyr Ile Arg Leu Ser Cys Asp Thr Asp Ala Glu Ile Leu Tyr
 65 70 75 80
 Glu Leu Leu Thr Gln His Trp His Leu Lys Thr Pro Asn Leu Val Ile
 85 90 95
 Ser Val Thr Gly Gly Ala Lys Asn Phe Ala Leu Lys Pro Arg Met Arg
 100 105 110
 Lys Ile Phe Ser Arg Leu Ile Tyr Ile Ala Gln Ser Lys Gly Ala Trp
 115 120 125
 Ile Leu Thr Gly Gly Thr His Tyr Gly Leu Met Lys Tyr Ile Gly Glu
 130 135 140
 Val Val Arg Asp Asn Thr Ile Ser Arg Ser Ser Glu Glu Asn Ile Val
 145 150 155 160
 Ala Ile Gly Ile Ala Ala Trp Gly Met Val Ser Asn Arg Asp Thr Leu
 165 170 175
 Ile Arg Asn Cys Asp Ala Glu Val Pro Val Gly Gln Glu Glu Val Cys
 180 185 190

<210> 15
 <211> 1095
 <212> PRT
 <213> Homo sapiens

<400> 15
 Met Arg Asn Arg Arg Asn Asp Thr Leu Asp Ser Thr Arg Thr Leu Tyr
 1 5 10 15
 Ser Ser Ala Ser Arg Ser Thr Asp Leu Ser Tyr Ser Glu Ser Asp Leu
 20 25 30
 Val Asn Phe Ile Gln Ala Asn Phe Lys Lys Arg Glu Cys Val Phe Phe
 35 40 45
 Thr Lys Asp Ser Lys Ala Thr Glu Asn Val Cys Lys Cys Gly Tyr Ala
 50 55 60
 Gln Ser Gln His Met Glu Gly Thr Gln Ile Asn Gln Ser Glu Lys Trp
 65 70 75 80

- 20 -

Asn Tyr Lys Lys His Thr Lys Glu Phe Pro Thr Asp Ala Phe Gly Asp
 85 90 95
 Ile Gln Phe Glu Thr Leu Gly Lys Lys Gly Lys Tyr Ile Arg Leu Ser
 100 105 110
 Cys Asp Thr Asp Ala Glu Ile Leu Tyr Glu Leu Leu Thr Gln His Trp
 115 120 125
 His Leu Lys Thr Pro Asn Leu Val Ile Ser Val Thr Gly Gly Ala Lys
 130 135 140
 Asn Phe Ala Leu Lys Pro Arg Met Arg Lys Ile Phe Ser Arg Leu Ile
 145 150 155 160
 Tyr Ile Ala Gln Ser Lys Gly Ala Trp Ile Leu Thr Gly Gly Thr His
 165 170 175
 Tyr Gly Leu Thr Lys Tyr Ile Gly Glu Val Val Arg Asp Asn Thr Ile
 180 185 190
 Ser Arg Ser Ser Glu Glu Asn Ile Val Ala Ile Gly Ile Ala Ala Trp
 195 200 205
 Gly Met Val Ser Asn Arg Asp Thr Leu Ile Arg Asn Cys Asp Ala Glu
 210 215 220
 Gly Tyr Phe Leu Ala Gln Tyr Leu Met Asp Asp Phe Thr Arg Asp Pro
 225 230 235 240
 Leu Tyr Ile Leu Asp Asn Asn His Thr His Leu Leu Leu Val Asp Asn
 245 250 255
 Gly Cys His Gly His Pro Thr Val Glu Ala Lys Leu Arg Asn Gln Leu
 260 265 270
 Glu Lys His Ile Ser Glu Arg Thr Ile Gln Asp Ser Asn Tyr Gly Gly
 275 280 285
 Lys Ile Pro Ile Val Cys Phe Ala Gln Gly Gly Gly Lys Glu Thr Leu
 290 295 300
 Lys Ala Ile Asn Thr Ser Ile Lys Asn Lys Ile Pro Cys Val Val Val
 305 310 315 320
 Glu Gly Ser Gly Arg Ile Ala Asp Val Ile Ala Ser Leu Val Glu Val
 325 330 335
 Glu Asp Ala Pro Thr Ser Ser Ala Val Lys Glu Lys Leu Val Arg Phe
 340 345 350
 Leu Pro Arg Thr Val Ser Arg Leu Ser Glu Glu Glu Thr Glu Ser Trp
 355 360 365
 Ile Lys Trp Leu Lys Glu Ile Leu Glu Cys Ser His Leu Leu Thr Val
 370 375 380
 Ile Lys Met Glu Glu Ala Gly Asp Glu Ile Val Ser Asn Ala Ile Ser
 385 390 395 400
 Tyr Ala Leu Tyr Lys Ala Phe Ser Thr Ser Glu Gln Asp Lys Asp Asn
 405 410 415
 Trp Asn Gly Gln Leu Lys Leu Leu Leu Glu Trp Asn Gln Leu Asp Leu
 420 425 430
 Ala Asn Asp Glu Ile Phe Thr Asn Asp Arg Arg Trp Glu Ser Ala Asp
 435 440 445

- 21 -

Leu Gln Glu Val Met Phe Thr Ala Leu Ile Lys Asp Arg Pro Lys Phe
 450 455 460
 Val Arg Leu Phe Leu Glu Asn Gly Leu Asn Leu Arg Lys Phe Leu Thr
 465 470 475 480
 His Asp Val Leu Thr Glu Leu Phe Ser Asn His Phe Ser Thr Leu Val
 485 490 495
 Tyr Arg Asn Leu Gln Ile Ala Lys Asn Ser Tyr Asn Asp Ala Leu Leu
 500 505 510
 Thr Phe Val Trp Lys Leu Val Ala Asn Phe Arg Arg Gly Phe Arg Lys
 515 520 525
 Glu Asp Arg Asn Gly Arg Asp Glu Met Asp Ile Glu Leu His Asp Val
 530 535 540
 Ser Pro Ile Thr Arg His Pro Leu Gln Ala Leu Phe Ile Trp Ala Ile
 545 550 555 560
 Leu Gln Asn Lys Lys Glu Leu Ser Lys Val Ile Trp Glu Gln Thr Arg
 565 570 575
 Gly Cys Thr Leu Ala Ala Leu Gly Ala Ser Lys Leu Leu Lys Thr Leu
 580 585 590
 Ala Lys Val Lys Asn Asp Ile Asn Ala Ala Gly Glu Ser Glu Glu Leu
 595 600 605
 Ala Asn Glu Tyr Glu Thr Arg Ala Val Glu Leu Phe Thr Glu Cys Tyr
 610 615 620
 Ser Ser Asp Glu Asp Leu Ala Glu Gln Leu Leu Val Tyr Ser Cys Glu
 625 630 635 640
 Ala Trp Gly Gly Ser Asn Cys Leu Glu Leu Ala Val Glu Ala Thr Asp
 645 650 655
 Gln His Phe Thr Ala Gln Pro Gly Val Gln Asn Phe Leu Ser Lys Gln
 660 665 670
 Trp Tyr Gly Glu Ile Ser Arg Asp Thr Lys Asn Trp Lys Ile Ile Leu
 675 680 685
 Cys Leu Phe Ile Ile Pro Leu Val Gly Cys Gly Phe Val Ser Phe Arg
 690 695 700
 Lys Lys Pro Val Asp Lys His Lys Lys Leu Leu Trp Tyr Tyr Val Ala
 705 710 715 720
 Phe Phe Thr Ser Pro Phe Val Val Phe Ser Trp Asn Val Val Phe Tyr
 725 730 735
 Ile Ala Phe Leu Leu Leu Phe Ala Tyr Val Leu Leu Met Asp Phe His
 740 745 750
 Ser Val Pro His Pro Pro Glu Leu Val Leu Tyr Ser Leu Val Phe Val
 755 760 765
 Leu Phe Cys Asp Glu Val Arg Gln Trp Tyr Val Asn Gly Val Asn Tyr
 770 775 780
 Phe Thr Asp Leu Trp Asn Val Met Asp Thr Leu Gly Leu Phe Tyr Phe
 785 790 795 800

- 22 -

| | | | | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|------|------|------|
| Ile | Ala | Gly | Ile | Val | Phe | Arg | Leu | His | Ser | Ser | Asn | Lys | Ser | Ser | Leu | 805 | 810 | 815 |
| Tyr | Ser | Gly | Arg | Val | Ile | Phe | Cys | Leu | Asp | Tyr | Ile | Ile | Phe | Thr | Leu | 820 | 825 | 830 |
| Arg | Leu | Ile | His | Ile | Phe | Thr | Val | Ser | Arg | Asn | Leu | Gly | Pro | Lys | Ile | 835 | 840 | 845 |
| Ile | Met | Leu | Gln | Arg | Met | Leu | Ile | Asp | Val | Phe | Phe | Phe | Leu | Phe | Leu | 850 | 855 | 860 |
| Phe | Ala | Val | Trp | Met | Val | Ala | Phe | Gly | Val | Ala | Arg | Gln | Gly | Ile | Leu | 865 | 870 | 875 |
| Arg | Gln | Asn | Glu | Gln | Arg | Trp | Arg | Trp | Ile | Phe | Arg | Ser | Val | Ile | Tyr | 885 | 890 | 895 |
| Glu | Pro | Tyr | Leu | Ala | Met | Phe | Gly | Gln | Val | Pro | Ser | Asp | Val | Asp | Gly | 900 | 905 | 910 |
| Thr | Thr | Tyr | Asp | Phe | Ala | His | Cys | Thr | Phe | Thr | Gly | Asn | Glu | Ser | Lys | 915 | 920 | 925 |
| Pro | Leu | Cys | Val | Glu | Leu | Asp | Glu | His | Asn | Leu | Pro | Arg | Phe | Pro | Glu | 930 | 935 | 940 |
| Trp | Ile | Thr | Ile | Pro | Leu | Val | Cys | Ile | Tyr | Met | Leu | Ser | Thr | Asn | Ile | 945 | 950 | 955 |
| Leu | Leu | Val | Asn | Leu | Leu | Val | Ala | Met | Phe | Gly | Tyr | Thr | Val | Gly | Thr | 965 | 970 | 975 |
| Val | Gln | Glu | Asn | Asn | Asp | Gln | Val | Trp | Lys | Phe | Gln | Arg | Tyr | Phe | Leu | 980 | 985 | 990 |
| Val | Gln | Glu | Tyr | Cys | Ser | Arg | Leu | Asn | Ile | Pro | Phe | Pro | Phe | Ile | Val | 995 | 1000 | 1005 |
| Phe | Ala | Tyr | Phe | Tyr | Met | Val | Val | Lys | Lys | Cys | Phe | Lys | Cys | Cys | Cys | 1010 | 1015 | 1020 |
| Lys | Glu | Lys | Asn | Met | Glu | Ser | Ser | Val | Cys | Cys | Phe | Lys | Asn | Glu | Asp | 1025 | 1030 | 1035 |
| Asn | Glu | Thr | Leu | Ala | Trp | Glu | Gly | Val | Met | Lys | Glu | Asn | Tyr | Leu | Val | 1045 | 1050 | 1055 |
| Lys | Ile | Asn | Thr | Lys | Ala | Asn | Asp | Thr | Ser | Glu | Glu | Met | Arg | His | Arg | 1060 | 1065 | 1070 |
| Phe | Arg | Gln | Leu | Asp | Thr | Lys | Leu | Asn | Asp | Leu | Lys | Gly | Leu | Leu | Lys | 1075 | 1080 | 1085 |
| Glu | Ile | Ala | Asn | Lys | Ile | Lys | | | | | | | | | | 1090 | 1095 | |

```
<210> 16
<211> 1095
<212> PRT
<213> Homo sapiens
```

<400> 16
Met Arg Asn Arg Arg Asn Asp Thr Leu Asp Ser Thr Arg Thr Leu Tyr
1 5 10 15

- 23 -

Ser Ser Ala Ser Arg Ser Thr Asp Leu Ser Tyr Ser Glu Ser Asp Leu
 20 25 30
 Val Asn Phe Ile Gln Ala Asn Phe Lys Lys Arg Glu Cys Val Phe Phe
 35 40 45
 Thr Lys Asp Ser Lys Ala Thr Glu Asn Val Cys Lys Cys Gly Tyr Ala
 50 55 60
 Gln Ser Gln His Met Glu Gly Thr Gln Ile Asn Gln Ser Glu Lys Trp
 65 70 75 80
 Asn Tyr Lys Lys His Thr Lys Glu Phe Pro Thr Asp Ala Phe Gly Asp
 85 90 95
 Ile Gln Phe Glu Thr Leu Gly Lys Lys Gly Lys Tyr Ile Arg Leu Ser
 100 105 110
 Cys Asp Thr Asp Ala Glu Ile Leu Tyr Glu Leu Leu Thr Gln His Trp
 115 120 125
 His Leu Lys Thr Pro Asn Leu Val Ile Ser Val Thr Gly Gly Ala Lys
 130 135 140
 Asn Phe Ala Leu Lys Pro Arg Met Arg Lys Ile Phe Ser Arg Leu Ile
 145 150 155 160
 Tyr Ile Ala Gln Ser Lys Gly Ala Trp Ile Leu Thr Gly Gly Thr His
 165 170 175
 Tyr Gly Leu Met Lys Tyr Ile Gly Glu Val Val Arg Asp Asn Thr Ile
 180 185 190
 Ser Arg Ser Ser Glu Glu Asn Ile Val Ala Ile Gly Ile Ala Ala Trp
 195 200 205
 Gly Met Val Ser Asn Arg Asp Thr Leu Ile Arg Asn Cys Asp Ala Glu
 210 215 220
 Gly Tyr Phe Leu Ala Gln Tyr Leu Met Asp Asp Phe Thr Arg Asp Pro
 225 230 235 240
 Leu Tyr Ile Leu Asp Asn Asn His Thr His Leu Leu Leu Val Asp Asn
 245 250 255
 Gly Cys His Gly His Pro Thr Val Glu Ala Lys Leu Arg Asn Gln Leu
 260 265 270
 Glu Lys Tyr Ile Ser Glu Arg Thr Ile Gln Asp Ser Asn Tyr Gly Gly
 275 280 285
 Lys Ile Pro Ile Val Cys Phe Ala Gln Gly Gly Gly Lys Glu Thr Leu
 290 295 300
 Lys Ala Ile Asn Thr Ser Ile Lys Asn Lys Ile Pro Cys Val Val Val
 305 310 315 320
 Glu Gly Ser Gly Gln Ile Ala Asp Val Ile Ala Ser Leu Val Glu Val
 325 330 335
 Glu Asp Ala Leu Thr Ser Ser Ala Val Lys Glu Lys Leu Val Arg Phe
 340 345 350
 Leu Pro Arg Thr Val Ser Arg Leu Pro Glu Glu Glu Thr Glu Ser Trp
 355 360 365
 Ile Lys Trp Leu Lys Glu Ile Leu Glu Cys Ser His Leu Leu Thr Val
 370 375 380

- 24 -

Ile Lys Met Glu Glu Ala Gly Asp Glu Ile Val Ser Asn Ala Ile Ser
 385 390 395 400
 Tyr Ala Leu Tyr Lys Ala Phe Ser Thr Ser Glu Gln Asp Lys Asp Asn
 405 410 415
 Trp Asn Gly Gln Leu Lys Leu Leu Leu Glu Trp Asn Gln Leu Asp Leu
 420 425 430
 Ala Asn Asp Glu Ile Phe Thr Asn Asp Arg Arg Trp Glu Ser Ala Asp
 435 440 445
 Leu Gln Glu Val Met Phe Thr Ala Leu Ile Lys Asp Arg Pro Lys Phe
 450 455 460
 Val Arg Leu Phe Leu Glu Asn Gly Leu Asn Leu Arg Lys Phe Leu Thr
 465 470 475 480
 His Asp Val Leu Thr Glu Leu Phe Ser Asn His Phe Ser Thr Leu Val
 485 490 495
 Tyr Arg Asn Leu Gln Ile Ala Lys Asn Ser Tyr Asn Asp Ala Leu Leu
 500 505 510
 Thr Phe Val Trp Lys Leu Val Ala Asn Phe Arg Arg Gly Phe Arg Lys
 515 520 525
 Glu Asp Arg Asn Gly Arg Asp Glu Met Asp Ile Glu Leu His Asp Val
 530 535 540
 Ser Pro Ile Thr Arg His Pro Leu Gln Ala Leu Phe Ile Trp Ala Ile
 545 550 555 560
 Leu Gln Asn Lys Lys Glu Leu Ser Lys Val Ile Trp Glu Gln Thr Arg
 565 570 575
 Gly Cys Thr Leu Ala Ala Leu Gly Ala Ser Lys Leu Leu Lys Thr Leu
 580 585 590
 Ala Lys Val Lys Asn Asp Ile Asn Ala Ala Gly Glu Ser Glu Glu Leu
 595 600 605
 Ala Asn Glu Tyr Glu Thr Arg Ala Val Glu Leu Phe Thr Glu Cys Tyr
 610 615 620
 Ser Ser Asp Glu Asp Leu Ala Glu Gln Leu Leu Val Tyr Ser Cys Glu
 625 630 635 640
 Ala Trp Gly Gly Ser Asn Cys Leu Glu Leu Ala Val Glu Ala Thr Asp
 645 650 655
 Gln His Phe Ile Ala Gln Pro Gly Val Gln Asn Phe Leu Ser Lys Gln
 660 665 670
 Trp Tyr Gly Glu Ile Ser Arg Asp Thr Lys Asn Trp Lys Ile Ile Leu
 675 680 685
 Cys Leu Phe Ile Ile Pro Leu Val Gly Cys Gly Phe Val Ser Phe Arg
 690 695 700
 Lys Lys Pro Val Asp Lys His Lys Lys Leu Leu Trp Tyr Tyr Val Ala
 705 710 715 720
 Phe Phe Thr Ser Pro Phe Val Val Phe Ser Trp Asn Val Val Phe Tyr
 725 730 735

- 25 -

Ile Ala Phe Leu Leu Leu Phe Ala Tyr Val Leu Leu Met Asp Phe His
 740 745 750
 Ser Val Pro His Pro Pro Glu Leu Val Leu Tyr Ser Leu Val Phe Val
 755 760 765
 Leu Phe Cys Asp Glu Val Arg Gln Trp Tyr Val Asn Gly Val Asn Tyr
 770 775 780
 Phe Thr Asp Leu Trp Asn Val Met Asp Thr Leu Gly Leu Phe Tyr Phe
 785 790 795 800
 Ile Ala Gly Ile Val Phe Arg Leu His Ser Ser Asn Lys Ser Ser Leu
 805 810 815
 Tyr Ser Gly Arg Val Ile Phe Cys Leu Asp Tyr Ile Ile Phe Thr Leu
 820 825 830
 Arg Leu Ile His Ile Phe Thr Val Ser Arg Asn Leu Gly Pro Lys Ile
 835 840 845
 Ile Met Leu Gln Arg Met Leu Ile Asp Val Phe Phe Phe Leu Phe Leu
 850 855 860
 Phe Ala Xaa Trp Met Val Ala Phe Gly Val Ala Arg Gln Gly Ile Leu
 865 870 875 880
 Arg Gln Asn Glu Gln Arg Trp Arg Trp Ile Phe Arg Ser Val Ile Tyr
 885 890 895
 Glu Pro Tyr Leu Ala Met Phe Gly Gln Val Pro Ser Asp Val Asp Gly
 900 905 910
 Thr Thr Tyr Asp Phe Ala His Cys Thr Phe Thr Gly Asn Glu Ser Lys
 915 920 925
 Pro Leu Cys Val Glu Leu Asp Glu His Asn Leu Pro Arg Phe Pro Glu
 930 935 940
 Trp Ile Thr Ile Pro Leu Val Cys Ile Tyr Met Leu Ser Thr Asn Ile
 945 950 955 960
 Leu Leu Val Asn Leu Leu Val Ala Met Phe Gly Tyr Thr Val Gly Thr
 965 970 975
 Val Gln Glu Asn Asn Asp Gln Val Trp Lys Phe Gln Arg Tyr Phe Leu
 980 985 990
 Val Gln Glu Tyr Cys Ser Arg Leu Asn Ile Pro Phe Pro Phe Ile Val
 995 1000 1005
 Phe Ala Tyr Phe Tyr Met Val Val Lys Lys Cys Phe Lys Cys Cys Cys
 1010 1015 1020
 Lys Glu Lys Asn Met Glu Ser Ser Val Cys Cys Phe Lys Asn Glu Asp
 1025 1030 1035 1040
 Asn Glu Thr Leu Ala Trp Glu Gly Val Met Lys Glu Asn Tyr Leu Val
 1045 1050 1055
 Lys Ile Asn Thr Lys Ala Asn Asp Thr Ser Glu Glu Met Arg His Arg
 1060 1065 1070
 Phe Arg Gln Leu Asp Thr Lys Leu Asn Asp Leu Lys Gly Leu Leu Lys
 1075 1080 1085
 Glu Ile Ala Asn Lys Ile Lys
 1090 1095

- 26 -

<210> 17
 <211> 652
 <212> PRT
 <213> Homo sapiens

<400> 17
 Met Arg Asn Arg Arg Asn Asp Thr Leu Asp Ser Thr Arg Thr Leu Tyr
 1 5 10 15
 Ser Ser Ala Ser Arg Ser Thr Asp Leu Ser Tyr Ser Glu Ser Asp Leu
 20 25 30
 Val Asn Phe Ile Gln Ala Asn Phe Lys Lys Arg Glu Cys Val Phe Phe
 35 40 45
 Thr Lys Asp Ser Lys Ala Thr Glu Asn Val Cys Lys Cys Gly Tyr Ala
 50 55 60
 Gln Ser Gln His Met Glu Gly Thr Gln Ile Asn Gln Ser Glu Lys Trp
 65 70 75 80
 Asn Tyr Lys Lys His Thr Lys Glu Phe Pro Thr Asp Ala Phe Gly Asp
 85 90 95
 Ile Gln Phe Glu Thr Leu Gly Lys Lys Gly Lys Tyr Ile Arg Leu Ser
 100 105 110
 Cys Asp Thr Asp Ala Glu Ile Leu Tyr Glu Leu Leu Thr Gln His Trp
 115 120 125
 His Leu Lys Thr Pro Asn Leu Val Ile Ser Val Thr Gly Gly Ala Lys
 130 135 140
 Asn Phe Ala Leu Lys Pro Arg Met Arg Lys Ile Phe Ser Arg Leu Ile
 145 150 155 160
 Tyr Ile Ala Gln Ser Lys Gly Ala Trp Ile Leu Thr Gly Gly Thr His
 165 170 175
 Tyr Gly Leu Met Lys Tyr Ile Gly Glu Val Val Arg Asp Asn Thr Ile
 180 185 190
 Ser Arg Ser Ser Glu Glu Asn Ile Val Ala Ile Gly Ile Ala Ala Trp
 195 200 205
 Gly Met Val Ser Asn Arg Asp Thr Leu Ile Arg Asn Cys Asp Ala Glu
 210 215 220
 Gly Tyr Phe Leu Ala Gln Tyr Leu Met Asp Asp Phe Thr Arg Asp Pro
 225 230 235 240
 Leu Tyr Ile Leu Asp Asn Asn His Thr His Leu Leu Leu Val Asp Asn
 245 250 255
 Gly Cys His Gly His Pro Thr Val Glu Ala Lys Leu Arg Asn Gln Leu
 260 265 270
 Glu Lys Tyr Ile Ser Glu Arg Thr Ile Gln Asp Ser Asn Tyr Gly Gly
 275 280 285
 Lys Ile Pro Ile Val Cys Phe Ala Gln Gly Gly Gly Lys Glu Thr Leu
 290 295 300
 Lys Ala Ile Asn Thr Ser Ile Lys Asn Lys Ile Pro Cys Val Val Val
 305 310 315 320

- 27 -

Glu Gly Ser Gly Gln Ile Ala Asp Val Ile Ala Ser Leu Val Glu Val
 325 330 335
 Glu Asp Ala Leu Thr Ser Ser Ala Val Lys Glu Lys Leu Val Arg Phe
 340 345 350
 Leu Pro Arg Thr Val Ser Arg Leu Pro Glu Glu Glu Thr Glu Ser Trp
 355 360 365
 Ile Lys Trp Leu Lys Glu Ile Leu Glu Cys Ser His Leu Leu Thr Val
 370 375 380
 Ile Lys Met Glu Glu Ala Gly Asp Glu Ile Val Ser Asn Ala Ile Ser
 385 390 395 400
 Tyr Ala Leu Tyr Lys Ala Phe Ser Thr Ser Glu Gln Asp Lys Asp Asn
 405 410 415
 Trp Asn Gly Gln Leu Lys Leu Leu Leu Glu Trp Asn Gln Leu Asp Leu
 420 425 430
 Ala Asn Asp Glu Ile Phe Thr Asn Asp Arg Arg Trp Glu Ser Ala Asp
 435 440 445
 Leu Gln Glu Val Met Phe Thr Ala Leu Ile Lys Asp Arg Pro Lys Phe
 450 455 460
 Val Arg Leu Phe Leu Glu Asn Gly Leu Asn Leu Arg Lys Phe Leu Thr
 465 470 475 480
 His Asp Val Leu Thr Glu Leu Phe Ser Asn His Phe Ser Thr Leu Val
 485 490 495
 Tyr Arg Asn Leu Gln Ile Ala Lys Asn Ser Tyr Asn Asp Ala Leu Leu
 500 505 510
 Thr Phe Val Trp Lys Leu Val Ala Asn Phe Arg Arg Gly Phe Arg Lys
 515 520 525
 Glu Asp Arg Asn Gly Arg Asp Glu Met Asp Ile Glu Leu His Asp Val
 530 535 540
 Ser Pro Ile Thr Arg His Pro Leu Gln Ala Leu Phe Ile Trp Ala Ile
 545 550 555 560
 Leu Gln Asn Lys Lys Glu Leu Ser Lys Val Ile Trp Glu Gln Thr Arg
 565 570 575
 Gly Cys Thr Leu Ala Ala Leu Gly Ala Ser Lys Leu Leu Lys Thr Leu
 580 585 590
 Ala Lys Val Lys Asn Asp Ile Asn Ala Ala Gly Glu Ser Glu Glu Leu
 595 600 605
 Ala Asn Glu Tyr Glu Thr Arg Ala Val Glu Leu Phe Thr Glu Cys Tyr
 610 615 620
 Ser Ser Asp Glu Asp Leu Ala Glu Gln Leu Leu Val Tyr Ser Cys Glu
 625 630 635 640
 Ala Trp Gly Gly Leu Glu His His His His His His
 645 650

<210> 18
 <211> 1095
 <212> PRT
 <213> Homo sapiens

- 28 -

<400> 18

Met Arg Asn Arg Arg Asn Asp Thr Leu Asp Ser Thr Arg Thr Leu Tyr
 1 5 10 15
 Ser Ser Ala Ser Arg Ser Thr Asp Leu Ser Tyr Ser Glu Ser Asp Leu
 20 25 30
 Val Asn Phe Ile Gln Ala Asn Phe Lys Lys Arg Glu Cys Val Phe Phe
 35 40 45
 Thr Lys Asp Ser Lys Ala Thr Glu Asn Val Cys Lys Cys Gly Tyr Ala
 50 55 60
 Gln Ser Gln His Met Glu Gly Thr Gln Ile Asn Gln Ser Glu Lys Trp
 65 70 75 80
 Asn Tyr Lys Lys His Thr Lys Glu Phe Pro Thr Asp Ala Phe Gly Asp
 85 90 95
 Ile Gln Phe Glu Thr Leu Gly Lys Lys Gly Lys Tyr Ile Arg Leu Ser
 100 105 110
 Cys Asp Thr Asp Ala Glu Ile Leu Tyr Glu Leu Leu Thr Gln His Trp
 115 120 125
 His Leu Lys Thr Pro Asn Leu Val Ile Ser Val Thr Gly Gly Ala Lys
 130 135 140
 Asn Phe Ala Leu Lys Pro Arg Met Arg Lys Ile Phe Ser Arg Leu Ile
 145 150 155 160
 Tyr Ile Ala Gln Ser Lys Gly Ala Trp Ile Leu Thr Gly Gly Thr His
 165 170 175
 Tyr Gly Leu Met Lys Tyr Ile Gly Glu Val Val Arg Asp Asn Thr Ile
 180 185 190
 Ser Arg Ser Ser Glu Glu Asn Ile Val Ala Ile Gly Ile Ala Ala Trp
 195 200 205
 Gly Met Val Ser Asn Arg Asp Thr Leu Ile Arg Asn Cys Asp Ala Glu
 210 215 220
 Gly Tyr Phe Leu Ala Gln Tyr Leu Met Asp Asp Phe Thr Arg Asp Pro
 225 230 235 240
 Leu Tyr Ile Leu Asp Asn Asn His Thr His Leu Leu Leu Val Asp Asn
 245 250 255
 Gly Cys His Gly His Pro Thr Val Glu Ala Lys Leu Arg Asn Gln Leu
 260 265 270
 Glu Lys Tyr Ile Ser Glu Arg Thr Ile Gln Asp Ser Asn Tyr Gly Gly
 275 280 285
 Lys Ile Pro Ile Val Cys Phe Ala Gln Gly Gly Gly Lys Glu Thr Leu
 290 295 300
 Lys Ala Ile Asn Thr Ser Ile Lys Asn Lys Ile Pro Cys Val Val Val
 305 310 315 320
 Glu Gly Ser Gly Gln Ile Ala Asp Val Ile Ala Ser Leu Val Glu Val
 325 330 335
 Glu Asp Ala Leu Thr Ser Ser Ala Val Lys Glu Lys Leu Val Arg Phe
 340 345 350

Leu Pro Arg Thr Val Ser Arg Leu Pro Glu Glu Glu Thr Glu Ser Trp
 355 360 365
 Ile Lys Trp Leu Lys Glu Ile Leu Glu Cys Ser His Leu Leu Thr Val
 370 375 380
 Ile Lys Met Glu Glu Ala Gly Asp Glu Ile Val Ser Asn Ala Ile Ser
 385 390 395 400
 Tyr Ala Leu Tyr Lys Ala Phe Ser Thr Ser Glu Gln Asp Lys Asp Asn
 405 410 415
 Trp Asn Gly Gln Leu Lys Leu Leu Leu Glu Trp Asn Gln Leu Asp Leu
 420 425 430
 Ala Asn Asp Glu Ile Phe Thr Asn Asp Arg Arg Trp Glu Ser Ala Asp
 435 440 445
 Leu Gln Glu Val Met Phe Thr Ala Leu Ile Lys Asp Arg Pro Lys Phe
 450 455 460
 Val Arg Leu Phe Leu Glu Asn Gly Leu Asn Leu Arg Lys Phe Leu Thr
 465 470 475 480
 His Asp Val Leu Thr Glu Leu Phe Ser Asn His Phe Ser Thr Leu Val
 485 490 495
 Tyr Arg Asn Leu Gln Ile Ala Lys Asn Ser Tyr Asn Asp Ala Leu Leu
 500 505 510
 Thr Phe Val Trp Lys Leu Val Ala Asn Phe Arg Arg Gly Phe Arg Lys
 515 520 525
 Glu Asp Arg Asn Gly Arg Asp Glu Met Asp Ile Glu Leu His Asp Val
 530 535 540
 Ser Pro Ile Thr Arg His Pro Leu Gln Ala Leu Phe Ile Trp Ala Ile
 545 550 555 560
 Leu Gln Asn Lys Lys Glu Leu Ser Lys Val Ile Trp Glu Gln Thr Arg
 565 570 575
 Gly Cys Thr Leu Ala Ala Leu Gly Ala Ser Lys Leu Leu Lys Thr Leu
 580 585 590
 Ala Lys Val Lys Asn Asp Ile Asn Ala Ala Gly Glu Ser Glu Glu Leu
 595 600 605
 Ala Asn Glu Tyr Glu Thr Arg Ala Val Glu Leu Phe Thr Glu Cys Tyr
 610 615 620
 Ser Ser Asp Glu Asp Leu Ala Glu Gln Leu Leu Val Tyr Ser Cys Glu
 625 630 635 640
 Ala Trp Gly Gly Ser Asn Cys Leu Glu Leu Ala Val Glu Ala Thr Asp
 645 650 655
 Gln His Phe Ile Ala Gln Pro Gly Val Gln Asn Phe Leu Ser Lys Gln
 660 665 670
 Trp Tyr Gly Glu Ile Ser Arg Asp Thr Lys Asn Trp Lys Ile Ile Leu
 675 680 685
 Cys Leu Phe Ile Ile Pro Leu Val Gly Cys Gly Phe Val Ser Phe Arg
 690 695 700
 Lys Lys Pro Val Asp Lys His Lys Lys Leu Leu Trp Tyr Tyr Val Ala
 705 710 715 720

- 30 -

Phe Phe Thr Ser Pro Phe Val Val Phe Ser Trp Asn Val Val Phe Tyr
 725 730 735
 Ile Ala Phe Leu Leu Leu Phe Ala Tyr Val Leu Leu Met Asp Phe His
 740 745 750
 Ser Val Pro His Pro Pro Glu Leu Val Leu Tyr Ser Leu Val Phe Val
 755 760 765
 Leu Phe Cys Asp Glu Val Arg Gln Trp Tyr Val Asn Gly Val Asn Tyr
 770 775 780
 Phe Thr Asp Leu Trp Asn Val Met Asp Thr Leu Gly Leu Phe Tyr Phe
 785 790 795 800
 Ile Ala Gly Ile Val Phe Arg Leu His Ser Ser Asn Lys Ser Ser Leu
 805 810 815
 Tyr Ser Gly Arg Val Ile Phe Cys Leu Asp Tyr Ile Ile Phe Thr Leu
 820 825 830
 Arg Leu Ile His Ile Phe Thr Val Ser Arg Asn Leu Gly Pro Lys Ile
 835 840 845
 Ile Met Leu Gln Arg Met Leu Ile Asp Val Phe Phe Phe Leu Phe Leu
 850 855 860
 Phe Ala Xaa Trp Met Val Ala Phe Gly Val Ala Arg Gln Gly Ile Leu
 865 870 875 880
 Arg Gln Asn Glu Gln Arg Trp Arg Trp Ile Phe Arg Ser Val Ile Tyr
 885 890 895
 Glu Pro Tyr Leu Ala Met Phe Gly Gln Val Pro Ser Asp Val Asp Gly
 900 905 910
 Thr Thr Tyr Asp Phe Ala His Cys Thr Phe Thr Gly Asn Glu Ser Lys
 915 920 925
 Pro Leu Cys Val Glu Leu Asp Glu His Asn Leu Pro Arg Phe Pro Glu
 930 935 940
 Trp Ile Thr Ile Pro Leu Val Cys Ile Tyr Met Leu Ser Thr Asn Ile
 945 950 955 960
 Leu Leu Val Asn Leu Leu Val Ala Met Phe Gly Tyr Thr Val Gly Thr
 965 970 975
 Val Gln Glu Asn Asn Asp Gln Val Trp Lys Phe Gln Arg Tyr Phe Leu
 980 985 990
 Val Gln Glu Tyr Cys Ser Arg Leu Asn Ile Pro Phe Pro Phe Ile Val
 995 1000 1005
 Phe Ala Tyr Phe Tyr Met Val Val Lys Lys Cys Phe Lys Cys Cys Cys
 1010 1015 1020
 Lys Glu Lys Asn Met Glu Ser Ser Val Cys Cys Phe Lys Asn Glu Asp
 1025 1030 1035 1040
 Asn Glu Thr Leu Ala Trp Glu Gly Val Met Lys Glu Asn Tyr Leu Val
 1045 1050 1055
 Lys Ile Asn Thr Lys Ala Asn Asp Thr Ser Glu Glu Met Arg His Arg
 1060 1065 1070

- 31 -

Phe Arg Gln Leu Asp Thr Lys Leu Asn Asp Leu Lys Gly Leu Leu Lys
 1075 1080 1085

Glu Ile Ala Asn Lys Ile Lys
 1090 1095

<210> 19
 <211> 652
 <212> PRT
 <213> Homo sapiens

<400> 19
 Met Arg Asn Arg Arg Asn Asp Thr Leu Asp Ser Thr Arg Thr Leu Tyr
 1 5 10 15

Ser Ser Ala Ser Arg Ser Thr Asp Leu Ser Tyr Ser Glu Ser Asp Leu
 20 25 30

Val Asn Phe Ile Gln Ala Asn Phe Lys Lys Arg Glu Cys Val Phe Phe
 35 40 45

Thr Lys Asp Ser Lys Ala Thr Glu Asn Val Cys Lys Cys Gly Tyr Ala
 50 55 60

Gln Ser Gln His Met Glu Gly Thr Gln Ile Asn Gln Ser Glu Lys Trp
 65 70 75 80

Asn Tyr Lys Lys His Thr Lys Glu Phe Pro Thr Asp Ala Phe Gly Asp
 85 90 95

Ile Gln Phe Glu Thr Leu Gly Lys Lys Gly Lys Tyr Ile Arg Leu Ser
 100 105 110

Cys Asp Thr Asp Ala Glu Ile Leu Tyr Glu Leu Leu Thr Gln His Trp
 115 120 125

His Leu Lys Thr Pro Asn Leu Val Ile Ser Val Thr Gly Gly Ala Lys
 130 135 140

Asn Phe Ala Leu Lys Pro Arg Met Arg Lys Ile Phe Ser Arg Leu Ile
 145 150 155 160

Tyr Ile Ala Gln Ser Lys Gly Ala Trp Ile Leu Thr Gly Gly Thr His
 165 170 175

Tyr Gly Leu Met Lys Tyr Ile Gly Glu Val Val Arg Asp Asn Thr Ile
 180 185 190

Ser Arg Ser Ser Glu Glu Asn Ile Val Ala Ile Gly Ile Ala Ala Trp
 195 200 205

Gly Met Val Ser Asn Arg Asp Thr Leu Ile Arg Asn Cys Asp Ala Glu
 210 215 220

Gly Tyr Phe Leu Ala Gln Tyr Leu Met Asp Asp Phe Thr Arg Asp Pro
 225 230 235 240

Leu Tyr Ile Leu Asp Asn Asn His Thr His Leu Leu Leu Val Asp Asn
 245 250 255

Gly Cys His Gly His Pro Thr Val Glu Ala Lys Leu Arg Asn Gln Leu
 260 265 270

Glu Lys Tyr Ile Ser Glu Arg Thr Ile Gln Asp Ser Asn Tyr Gly Gly
 275 280 285

- 32 -

Lys Ile Pro Ile Val Cys Phe Ala Gln Gly Gly Gly Lys Glu Thr Leu
 290 295 300
 Lys Ala Ile Asn Thr Ser Ile Lys Asn Lys Ile Pro Cys Val Val Val
 305 310 315 320
 Glu Gly Ser Gly Gln Ile Ala Asp Val Ile Ala Ser Leu Val Glu Val
 325 330 335
 Glu Asp Ala Leu Thr Ser Ser Ala Val Lys Glu Lys Leu Val Arg Phe
 340 345 350
 Leu Pro Arg Thr Val Ser Arg Leu Pro Glu Glu Glu Thr Glu Ser Trp
 355 360 365
 Ile Lys Trp Leu Lys Glu Ile Leu Glu Cys Ser His Leu Leu Thr Val
 370 375 380
 Ile Lys Met Glu Glu Ala Gly Asp Glu Ile Val Ser Asn Ala Ile Ser
 385 390 395 400
 Tyr Ala Leu Tyr Lys Ala Phe Ser Thr Ser Glu Gln Asp Lys Asp Asn
 405 410 415
 Trp Asn Gly Gln Leu Lys Leu Leu Leu Glu Trp Asn Gln Leu Asp Leu
 420 425 430
 Ala Asn Asp Glu Ile Phe Thr Asn Asp Arg Arg Trp Glu Ser Ala Asp
 435 440 445
 Leu Gln Glu Val Met Phe Thr Ala Leu Ile Lys Asp Arg Pro Lys Phe
 450 455 460
 Val Arg Leu Phe Leu Glu Asn Gly Leu Asn Leu Arg Lys Phe Leu Thr
 465 470 475 480
 His Asp Val Leu Thr Glu Leu Phe Ser Asn His Phe Ser Thr Leu Val
 485 490 495
 Tyr Arg Asn Leu Gln Ile Ala Lys Asn Ser Tyr Asn Asp Ala Leu Leu
 500 505 510
 Thr Phe Val Trp Lys Leu Val Ala Asn Phe Arg Arg Gly Phe Arg Lys
 515 520 525
 Glu Asp Arg Asn Gly Arg Asp Glu Met Asp Ile Glu Leu His Asp Val
 530 535 540
 Ser Pro Ile Thr Arg His Pro Leu Gln Ala Leu Phe Ile Trp Ala Ile
 545 550 555 560
 Leu Gln Asn Lys Lys Glu Leu Ser Lys Val Ile Trp Glu Gln Thr Arg
 565 570 575
 Gly Cys Thr Leu Ala Ala Leu Gly Ala Ser Lys Leu Leu Lys Thr Leu
 580 585 590
 Ala Lys Val Lys Asn Asp Ile Asn Ala Ala Gly Glu Ser Glu Glu Leu
 595 600 605
 Ala Asn Glu Tyr Glu Thr Arg Ala Val Glu Leu Phe Thr Glu Cys Tyr
 610 615 620
 Ser Ser Asp Glu Asp Leu Ala Glu Gln Leu Leu Val Tyr Ser Cys Glu
 625 630 635 640
 Ala Trp Gly Gly Leu Glu His His His His His His
 645 650

- 33 -

<210> 20
 <211> 1104
 <212> PRT
 <213> Homo sapiens

<400> 20
 Met Ser Phe Arg Ala Ala Arg Leu Ser Met Arg Asn Arg Arg Asn Asp
 1 5 10 15
 Thr Leu Asp Ser Thr Arg Thr Leu Tyr Ser Ser Ala Ser Arg Ser Thr
 20 25 30
 Asp Leu Ser Tyr Ser Glu Ser Asp Leu Val Asn Phe Ile Gln Ala Asn
 35 40 45
 Phe Lys Lys Arg Glu Cys Val Phe Phe Thr Lys Asp Ser Lys Ala Thr
 50 55 60
 Glu Asn Val Cys Lys Cys Gly Tyr Ala Gln Ser Gln His Met Glu Gly
 65 70 75 80
 Thr Gln Ile Asn Gln Ser Glu Lys Trp Asn Tyr Lys Lys His Thr Lys
 85 90 95
 Glu Phe Pro Thr Asp Ala Phe Gly Asp Ile Gln Phe Glu Thr Leu Gly
 100 105 110
 Lys Lys Gly Lys Tyr Ile Arg Leu Ser Cys Asp Thr Asp Ala Glu Ile
 115 120 125
 Leu Tyr Glu Leu Leu Thr Gln His Trp His Leu Lys Thr Pro Asn Leu
 130 135 140
 Val Ile Ser Val Thr Gly Gly Ala Lys Asn Phe Ala Leu Lys Pro Arg
 145 150 155 160
 Met Arg Lys Ile Phe Ser Arg Leu Ile Tyr Ile Ala Gln Ser Lys Gly
 165 170 175
 Ala Trp Ile Leu Thr Gly Gly Thr His Tyr Gly Leu Met Lys Tyr Ile
 180 185 190
 Gly Glu Val Val Arg Asp Asn Thr Ile Ser Arg Ser Ser Glu Glu Asn
 195 200 205
 Ile Val Ala Ile Gly Ile Ala Ala Trp Gly Met Val Ser Asn Arg Asp
 210 215 220
 Thr Leu Ile Arg Asn Cys Asp Ala Glu Gly Tyr Phe Leu Ala Gln Tyr
 225 230 235 240
 Leu Met Asp Asp Phe Thr Arg Asp Pro Leu Cys Ile Leu Asp Asn Asn
 245 250 255
 His Thr His Leu Leu Leu Val Asp Asn Gly Cys His Gly His Pro Thr
 260 265 270
 Val Glu Ala Lys Leu Arg Asn Gln Leu Glu Lys Tyr Ile Ser Glu Arg
 275 280 285
 Thr Ile Gln Asp Ser Asn Tyr Gly Gly Lys Ile Pro Ile Val Cys Phe
 290 295 300
 Ala Gln Gly Gly Gly Lys Glu Thr Leu Lys Ala Ile Asn Thr Ser Ile
 305 310 315 320

-34-

Lys Asn Lys Ile Pro Cys Val Val Val Glu Gly Ser Gly Gln Ile Ala
 325 330 335
 Asp Val Ile Ala Ser Leu Val Glu Val Glu Asp Ala Leu Thr Ser Ser
 340 345 350
 Ala Val Lys Glu Lys Leu Val Arg Phe Leu Pro Arg Thr Val Ser Arg
 355 360 365
 Leu Pro Glu Glu Glu Thr Glu Ser Trp Ile Lys Trp Leu Lys Glu Ile
 370 375 380
 Leu Glu Cys Ser His Leu Leu Thr Val Ile Lys Met Glu Glu Ala Gly
 385 390 395 400
 Asp Glu Ile Val Ser Asn Ala Ile Ser Tyr Ala Leu Tyr Lys Ala Phe
 405 410 415
 Ser Thr Ser Glu Gln Asp Lys Asp Asn Trp Asn Gly Gln Leu Lys Leu
 420 425 430
 Leu Leu Glu Trp Asn Gln Leu Asp Leu Ala Asn Asp Glu Ile Phe Thr
 435 440 445
 Asn Asp Arg Arg Trp Glu Ser Ala Asp Leu Gln Glu Val Met Phe Thr
 450 455 460
 Ala Leu Ile Lys Asp Arg Pro Lys Phe Val Arg Leu Phe Leu Glu Asn
 465 470 475 480
 Gly Leu Asn Leu Arg Lys Phe Leu Thr His Asp Val Leu Thr Glu Leu
 485 490 495
 Phe Ser Asn His Phe Ser Thr Leu Val Tyr Arg Asn Leu Gln Ile Ala
 500 505 510
 Lys Asn Ser Tyr Asn Asp Ala Leu Leu Thr Phe Val Trp Lys Leu Val
 515 520 525
 Ala Asn Phe Arg Arg Gly Phe Arg Lys Glu Asp Arg Asn Gly Arg Asp
 530 535 540
 Glu Met Asp Ile Glu Leu His Asp Val Ser Pro Ile Thr Arg His Pro
 545 550 555 560
 Leu Gln Ala Leu Phe Ile Trp Ala Ile Leu Gln Asn Lys Lys Glu Leu
 565 570 575
 Ser Lys Val Ile Trp Glu Gln Thr Arg Gly Cys Thr Leu Ala Ala Leu
 580 585 590
 Gly Ala Ser Lys Leu Leu Lys Thr Leu Ala Lys Val Lys Asn Asp Ile
 595 600 605
 Asn Ala Ala Gly Glu Ser Glu Glu Leu Ala Asn Glu Tyr Glu Thr Arg
 610 615 620
 Ala Val Glu Leu Phe Thr Glu Cys Tyr Ser Ser Asp Glu Asp Leu Ala
 625 630 635 640
 Glu Gln Leu Leu Val Tyr Ser Cys Glu Ala Trp Gly Gly Ser Asn Cys
 645 650 655
 Leu Glu Leu Ala Val Glu Ala Thr Asp Gln His Phe Ile Ala Gln Pro
 660 665 670
 Gly Val Gln Asn Phe Leu Ser Lys Gln Trp Tyr Gly Glu Ile Ser Arg
 675 680 685

Asp Thr Lys Asn Trp Lys Ile Ile Leu Cys Leu Phe Ile Ile Pro Leu
 690 695 700
 Val Gly Cys Gly Phe Val Ser Phe Arg Lys Lys Pro Val Asp Lys His
 705 710 715 720
 Lys Lys Leu Leu Trp Tyr Tyr Val Ala Phe Phe Thr Ser Pro Phe Val
 725 730 735
 Val Phe Ser Trp Asn Val Val Phe Tyr Ile Ala Phe Leu Leu Leu Phe
 740 745 750
 Ala Tyr Val Leu Leu Met Asp Phe His Ser Val Pro His Pro Pro Glu
 755 760 765
 Leu Val Leu Tyr Ser Leu Val Phe Val Leu Phe Cys Asp Glu Val Arg
 770 775 780
 Gln Trp Tyr Val Asn Gly Val Asn Tyr Phe Thr Asp Leu Trp Asn Val
 785 790 795 800
 Met Asp Thr Leu Gly Leu Phe Tyr Phe Ile Ala Gly Ile Val Phe Arg
 805 810 815
 Leu His Ser Ser Asn Lys Ser Ser Leu Tyr Ser Gly Arg Val Ile Phe
 820 825 830
 Cys Leu Asp Tyr Ile Ile Phe Thr Leu Arg Leu Ile His Ile Phe Thr
 835 840 845
 Val Ser Arg Asn Leu Gly Pro Lys Ile Ile Met Leu Gln Arg Met Leu
 850 855 860
 Ile Asp Val Phe Phe Phe Leu Phe Leu Phe Ala Val Trp Met Val Ala
 865 870 875 880
 Phe Gly Val Ala Arg Gln Gly Ile Leu Arg Gln Asn Glu Gln Arg Trp
 885 890 895
 Arg Trp Ile Phe Arg Ser Val Ile Tyr Glu Pro Tyr Leu Ala Met Phe
 900 905 910
 Gly Gln Val Pro Ser Asp Val Asp Gly Thr Thr Tyr Asp Phe Ala His
 915 920 925
 Cys Thr Phe Thr Gly Asn Glu Ser Lys Pro Leu Cys Val Glu Leu Asp
 930 935 940
 Glu His Asn Leu Pro Arg Phe Pro Glu Trp Ile Thr Ile Pro Leu Val
 945 950 955 960
 Cys Ile Tyr Met Leu Ser Thr Asn Ile Leu Leu Val Asn Leu Leu Val
 965 970 975
 Ala Met Phe Gly Tyr Thr Val Gly Thr Val Gln Glu Asn Asn Asp Gln
 980 985 990
 Val Trp Lys Phe Gln Arg Tyr Phe Leu Val Gln Glu Tyr Cys Ser Arg
 995 1000 1005
 Leu Asn Ile Pro Phe Pro Phe Ile Val Phe Ala Tyr Phe Tyr Met Val
 1010 1015 1020
 Val Lys Lys Cys Phe Lys Cys Cys Cys Lys Glu Lys Asn Met Glu Ser
 1025 1030 1035 1040

- 36 -

Ser Val Cys Cys Phe Lys Asn Glu Asp Asn Glu Thr Leu Ala Trp Glu
1045 1050 1055

Gly Val Met Lys Glu Asn Tyr Leu Val Lys Ile Asn Thr Lys Ala Asn
1060 1065 1070

Asp Thr Ser Glu Glu Met Arg His Arg Phe Arg Gln Leu Asp Thr Lys
1075 1080 1085

Leu Asn Asp Leu Lys Gly Leu Leu Lys Glu Ile Ala Asn Lys Ile Lys
1090 1095 1100

<210> 21
<211> 931
<212> PRT
<213> Homo sapiens

<400> 21
Met Val Gly Gly Cys Arg Trp Thr Glu Asp Val Glu Pro Ala Glu Val
1 5 10 15

Lys Glu Lys Met Ser Phe Arg Ala Ala Arg Leu Ser Met Arg Asn Arg
20 25 30

Arg Asn Asp Thr Leu Asp Ser Thr Arg Thr Leu Tyr Ser Ser Ala Ser
35 40 45

Arg Ser Thr Asp Leu Ser Tyr Ser Glu Ser Ala Ser Phe Tyr Ala Ala
50 55 60

Phe Arg Thr Gln Thr Cys Pro Ile Met Ala Ser Trp Asp Leu Val Asn
65 70 75 80

Phe Ile Gln Ala Asn Phe Lys Lys Arg Glu Cys Val Phe Phe Thr Lys
85 90 95

Asp Ser Lys Ala Thr Glu Asn Val Cys Lys Cys Gly Tyr Ala Gln Ser
100 105 110

Gln His Met Glu Gly Thr Gln Ile Asn Gln Ser Glu Lys Trp Asn Tyr
115 120 125

Lys Lys His Thr Lys Glu Phe Pro Thr Asp Ala Phe Gly Asp Ile Gln
130 135 140

Phe Glu Thr Leu Gly Lys Lys Gly Lys Tyr Ile Arg Leu Ser Cys Asp
145 150 155 160

Thr Asp Ala Glu Ile Leu Tyr Glu Leu Leu Thr Gln His Trp His Leu
165 170 175

Lys Thr Pro Asn Leu Val Ile Ser Val Thr Gly Gly Ala Lys Asn Phe
180 185 190

Ala Leu Lys Pro Arg Met Arg Lys Ile Phe Ser Arg Leu Ile Tyr Ile
195 200 205

Ala Gln Ser Lys Gly Ala Trp Ile Leu Thr Gly Gly Thr His Tyr Gly
210 215 220

Leu Met Lys Tyr Ile Gly Glu Val Val Arg Asp Asn Thr Ile Ser Arg
225 230 235 240

Ser Ser Glu Glu Asn Ile Val Ala Ile Gly Ile Ala Ala Trp Gly Met
 245 250 255
 Val Ser Asn Arg Asp Thr Leu Ile Arg Asn Cys Asp Ala Glu Gly Tyr
 260 265 270
 Phe Leu Ala Gln Tyr Leu Met Asp Asp Phe Thr Arg Asp Pro Leu Tyr
 275 280 285
 Ile Leu Asp Asn Asn His Thr His Leu Leu Leu Val Asp Asn Gly Cys
 290 295 300
 His Gly His Pro Thr Val Glu Ala Lys Leu Arg Asn Gln Leu Glu Lys
 305 310 315 320
 Tyr Ile Ser Glu Arg Thr Ile Gln Asp Ser Asn Tyr Gly Gly Lys Ile
 325 330 335
 Pro Ile Val Cys Phe Ala Gln Gly Gly Gly Lys Glu Thr Leu Lys Ala
 340 345 350
 Ile Asn Thr Ser Ile Lys Asn Lys Ile Pro Cys Val Val Val Glu Gly
 355 360 365
 Ser Gly Gln Ile Ala Asp Val Ile Ala Ser Leu Val Glu Val Glu Asp
 370 375 380
 Ala Leu Thr Ser Ser Ala Val Lys Glu Lys Leu Val Arg Phe Leu Pro
 385 390 395 400
 Arg Thr Val Ser Arg Leu Pro Glu Glu Glu Thr Glu Ser Trp Ile Lys
 405 410 415
 Trp Leu Lys Glu Ile Leu Glu Cys Ser His Leu Leu Thr Val Ile Lys
 420 425 430
 Met Glu Glu Ala Gly Asp Glu Ile Val Ser Asn Ala Ile Ser Tyr Ala
 435 440 445
 Leu Tyr Lys Ala Phe Ser Thr Ser Glu Gln Asp Lys Asp Asn Trp Asn
 450 455 460
 Gly Gln Leu Lys Leu Leu Leu Glu Trp Asn Gln Leu Asp Leu Ala Asn
 465 470 475 480
 Asp Glu Ile Phe Thr Asn Asp Arg Arg Trp Glu Lys Ser Lys Pro Arg
 485 490 495
 Leu Arg Asp Thr Ile Ile Gln Val Thr Trp Leu Glu Asn Gly Arg Ile
 500 505 510
 Lys Val Glu Ser Lys Asp Val Thr Asp Gly Lys Ala Ser Ser His Met
 515 520 525
 Leu Val Val Leu Lys Ser Ala Asp Leu Gln Glu Val Met Phe Thr Ala
 530 535 540
 Leu Ile Lys Asp Arg Pro Lys Phe Val Arg Leu Phe Leu Glu Asn Gly
 545 550 555 560
 Leu Asn Leu Arg Lys Phe Leu Thr His Asp Val Leu Thr Glu Leu Phe
 565 570 575
 Ser Asn His Phe Ser Thr Leu Val Tyr Arg Asn Leu Gln Ile Ala Lys
 580 585 590
 Asn Ser Tyr Asn Asp Ala Leu Leu Thr Phe Val Trp Lys Leu Val Ala
 595 600 605

Asn Phe Arg Arg Gly Phe Arg Lys Glu Asp Arg Asn Gly Arg Asp Glu
 610 615 620
 Met Asp Ile Glu Leu His Asp Val Ser Pro Ile Thr Arg His Pro Leu
 625 630 635 640
 Gln Ala Leu Phe Ile Trp Ala Ile Leu Gln Asn Lys Lys Glu Leu Ser
 645 650 655
 Lys Val Ile Trp Glu Gln Thr Arg Gly Cys Thr Leu Ala Ala Leu Gly
 660 665 670
 Ala Ser Lys Leu Leu Lys Thr Leu Ala Lys Val Lys Asn Asp Ile Asn
 675 680 685
 Ala Ala Gly Glu Ser Glu Glu Leu Ala Asn Glu Tyr Glu Thr Arg Ala
 690 695 700
 Val Gly Glu Ser Thr Val Trp Asn Ala Val Val Gly Ala Asp Leu Pro
 705 710 715 720
 Cys Gly Thr Asp Ile Ala Ser Gly Thr His Arg Pro Asp Gly Gly Glu
 725 730 735
 Leu Phe Thr Glu Cys Tyr Ser Ser Asp Glu Asp Leu Ala Glu Gln Leu
 740 745 750
 Leu Val Tyr Ser Cys Glu Ala Trp Gly Gly Ser Asn Cys Leu Glu Leu
 755 760 765
 Ala Val Glu Ala Thr Asp Gln His Phe Ile Ala Gln Pro Gly Val Gln
 770 775 780
 Asn Phe Leu Ser Lys Gln Trp Tyr Gly Glu Ile Ser Arg Asp Thr Lys
 785 790 795 800
 Asn Trp Lys Ile Ile Leu Cys Leu Phe Ile Ile Pro Leu Val Gly Cys
 805 810 815
 Gly Phe Val Ser Phe Arg Lys Lys Pro Val Asp Lys His Lys Lys Leu
 820 825 830
 Leu Trp Tyr Tyr Val Ala Phe Phe Thr Ser Pro Phe Val Val Phe Ser
 835 840 845
 Trp Asn Val Val Phe Tyr Ile Ala Phe Leu Leu Leu Phe Ala Tyr Val
 850 855 860
 Leu Leu Met Asp Phe His Ser Val Pro His Pro Pro Glu Leu Val Leu
 865 870 875 880
 Tyr Ser Leu Val Phe Val Leu Phe Cys Asp Glu Lys Arg Lys Thr Ala
 885 890 895
 Met Asp Gln Thr Asp Glu Asp Leu Phe Pro Tyr Gly Ala Phe Tyr Gln
 900 905 910
 Phe Leu Met Ile Ser Arg Ser Phe Arg Gly Glu Glu Met Ser Ile Gly
 915 920 925
 Lys Gln His
 930